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GenCore version 5.1.3 Copyright (c) 1993 - 2002 Compugen Ltd.
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OM nucleic - nucleic search, using sw model	Run on: November 19, 2002, 00:33:44; Search time 1026.35 Seconds (without alignments) 765.605 Million cell updates/sec	Title: US-09-829-004A-9 Perfect score: 27 Sequence: 1 aaacttcagtgtgtggacctccatgtt 27	Scoring table: IDENTITY_NUC Gapor 10.0 , Gapext 1.0	Searched: 2054640 segs, 14551402878 residues	Total number of hits satisfying chosen parameters: 4109280	Minimum DB seq length: 0 Maximum DB seq length: 2000000000	Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries
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Database :

GenEmbl:\*

1: 9b\_ba:\*
2: 9b\_htg:\*
3: 9b\_in:\*
3: 9b\_on:\*
3b\_ov:\*
9b\_pat:\*
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em\_htgo\_mus:\*
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Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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	27	100.	27		52		AX281	252 Sequence
71 6	77	100.0	737		HSPSA4		XISYA	A Human PSA
J 4	27	100.0	711		BD007601		BD0076	601 Method fo
Ŋ	27	100.0	870		HSA459783		AJ459	783 Homo sa
ωı	27	100.0	876		HSA459782		AJ459	782 Homo sapi
<b>~</b> 0	77	100.0	066		HSU1/040		U1/04	O Human prost
ာတ	27	100.0	1130		AF335477		AF335	477 Homo sapi
10	27	100.0	1350		AX200986		AX200	986 Sequence
11	27	100.0	1350		AX267642		AX267	642 Sequence
12	27	100.0	1415		•		M2189	5 Human prost
13	27	100.0	1446		HUMAPS		M2666	3 Homo sapien
14	27	100.0	1462		167864		16786	4 Sequence 14
15	77	100	1466		E32812		E3281	2 Primer DNA
170	7,0	200	1490		BC005307		2000	307 Homo sant
18	27	100.0	1603		HSA310938		AJ310	1938 Homo sapi
19	27	100.0	1654		HUMPAB		M2189	6 Human prost
20	27	100.0	1729		AR059540 .		AR059	540 Sequence
21	27	100.0	1729		AR082934		AR082	934 Sequence
22	27	100.0	1729		AX337560		AX337	550 Sequence
233	77	100.0	10.75		HSPSA NECOSATO		XU//3	10 Human mKNA
2.4 2.5	7 10	100	2106		AX106218		AX106	218 Seguence
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31	7 0	100.0	6153		HUMPSANTIG		M2454	3 Human prost
2 5 5	77	100.0	7130		A3/202		A5/20	2 sequence z
0 P	7,0	100	7130		HIMDSAA		M2727	733 Sequence
, c.	27	100.0	40458		AC011523		AC011	523 Homo sapi
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8	25.4	94.1			MMPROS		X7356	50 M.mulatta m
6	21.2	78.5	146276		AC090889		AC090	1889 Canis fam
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7 (	20.6	76.3	υ,		AC016/10		ACOL	omo sa
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0 0 4 4 4 5 5	200	74.1	20 20	ی م	119503		119503 Seq 167851 Sea	13 Sequence 1
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AX281252								
rocus	AX2		,		27 bp	DNA	linear	PAT 02-NOV-2001
DEFINITION	Sed		9 from E	Patent	W00176			
VERSION	AX2	AX281252.1	1 GI:1660850	9099	508			
KEYWORDS		4044		4				
SCURCE	syn	synthetic o	construct	ict.				
	art	ificia	l sequences	ces				

1 Pedyczak, A., Chong, P. and Sia, C.D. Immunogenic peptides derived from prostate-specific antigen (psa) and uses thereof Patent: WO 0176622-A 9 18-OCT-2001; REFERENCE AUTHORS TITLE

JOURNAL

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BD0075601.
BD007501.
BD0075601.
BD0075601.
BS Tindall,D.J., Young, C.Y.F., Mccormic,D.J., Klee,G.G., Saedi,M.S., Kundar,A., Tittenhouse,H.G. and Wolfert,R.L.
Brene: JP 2001503991-A.14 27-MAR-2001;
BATCH: JP 2001503991-A.14
BAYO. BCOUNATION FOR MEDICAL EDUCATION AND RESEARCH, HYBRITECH INC OS Unidentified
BN JP 2001503991-A.14
BN JP 200150991-A.14
BN MANAAD SAEED SAEDI,ABHAY KUMAR,HARRY G RITTENHOUSE, PI
BN MANAAD SAED SAEDI S
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1. .71
/organism='Unidentified'
Location/Qualifiers
                                                                                                                                                                                                  M Unknown.
Unclassified.
I (bases 1 to 711)
S Young, C.Y.F., Tindall, D.J. and Klee, G.G.
Method for detection of breast cancer
AL Patent: US 6235486-A 4 22-MAX-2001;
Location/Qualifiers
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Submitted (09-JAN-1989) Digby M.R., The Howard Florey Institute,
Nelbourne Dniversity, Parkville, Victoria 3052, Australia
2 (bases 1 to 281)
Digby,M., Zhang,X.Y. and Richards,R.I.
Human prostate specific antigen (PSA) gene: structure and linkage to the kallikrein-like gene, hGK-1
891886232
2467258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammaliai; Eutheria; Primates; Catarrhini; Hominidae; Homo.
Digby, M.R.
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100.0%; Score 27; DB 9; Length 281;
Best Local Similarity 100.0%; Pred. No. 0.011;
Matches 27; Conservative 0; Mismatches 0, Indels
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Human PSA gene for prostate specific antigen exon
X13943.1
X13943.1 GI:35728
Kallikrein; prostate specific antigen; PSA gene.
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1. 52
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Matches 27; Conservative 0; Mismatches 0;
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Aventis Pasteur Limited (CA)
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Gaps



Euteleostomi;

Craniata; Vertebrata; Eutele Catarrhini; Hominidae; Homo

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/translation="MWVPVVFLTLSVTWIGAAPLILSRIVGGWECEKHSQPWQVLVAS
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WTGGKSTCSGDSGGPLVCNGVLQGITSWGSEPCALPERPSLYTKVVHYRKWIKDTIVA
                                                                                                                                                           Heuze-Vourc'h,N. and Courty,Y. Complex alternative splicing of the hKLK3 gene coding for the tumour marker PSA (prostate-specific-antigen)
                                                                                                                                                                                                                                                                      Direct Submission
Submitted (03-MAY-2002) Courty Y., EMI-U 0010, Laboratoire Submitted (03-MAY-2002)
d'Enzynologie, Universite F. Rabelais, 2 bis byd Tonnelle, 37032 cedex, FRANCE alternative splice variant sequences: AJ459783, AJ459784.
Location/Qualifiers
1. 876
                                   AJ459782.1 GI:20520640 "... alternative splicing; KLK3 gene; prostate specific antigen
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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22. .684
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Courty, Y.
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                  AJ459782
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m HSA459782} 876 bp mRNA linear PRI 09-MAY-2002 Homo sapiens mRNA for prostate specific antigen (KLK3 gene), splice
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Homo sapiens mRNA for prostate specific antigen (KLK3 gene), splice
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Submitted (03-MAY-2002) Courty Y., EMI-U 0010, Laboratoire
d'Enzymologie, Universite F. Rabelais, 2 bis byd Tonnelle, Tours,
                                                                                                                                                                                                                                                                                                                                                             Heuze-Vourc'h,N. and Courty,Y.
Complex alternative splicing of the hKLK3 gene coding for the
tumour marker PSA (prostate-specific-antigen)
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alternative splicing; KLK3 gene; prostate specific antigen
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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/protein_id="CAD30845.1"
/db_xref="G1:20520643"
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 Mismatches
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/db_xref="taxon:9606"
/clone="YC140405-00"
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94. .804
/gene="KLK3"
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22. .807
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27; Conservative
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Courty, Y.
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A, complete cds.
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1 (bases 1 to 990)
Monne, M.W., Moreno, J.M., Mele, C.M., Mulholland, G.M. and
Gomella, L.G.
A Comparative Analysis of Prostate Specific Antigen Gene Sequence
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostom1;
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                                                                                               Length 876
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/product="prostate specific antigen"
a 235 c 231 g 190 t
                                                                                            Ouery Match
100.0%; Score 27; DB 9;
Best Local Similarity 100.0%; Pred. No. 0.012;
Matches 27; Conservative 0; Mismatches 0;
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Homo sapiens.

Homo sapiens.

Homo sapiens.

Homo sapiens.

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; CE | Chases 1 to 1130)

Baid, A., Mabjeesh, N., Azar, I., Biton, S., Engel, S., Bernstein, J., Lifschitz Mercer, B., Matzkin, H., Rotman, G., Toporik, A., Savitsky, K. and Mintz, L.

AL Singla alternative splicing within the human kallikrein genes KLK2 and KLK3 gives rise to novel prostate-specific proteins

AL Biol. Chem. 277 (20), 18084-18090 (2002)

E 22001265

ED 11834722

E Cases 1 to 1130)

Rad Mintz, L.

Bavid, A., Engel, S., Azar, I., Bernstein, J., Rotman, G., Savitsky, K.

Direct Submission

Submitted (11-JAN-2001) Compugen Ltd., 72 Pinchas Rosen, Tel Aviv
                                                                                            Homo sapiens prostate-specific antigen variant 1 mRNA, complete cds, alternatively spliced.
AF335477.1 GI:18478569
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/protein_id="AAL/3204.1"
/db_xref="G1:ld478570"
/translation="MWVPVVFLTLSVTWIGERGHGWGDAGEGASPDCOAEALSPPTOH
PSPDRELGSFLSLPAPLQAHTPSPSILCQSSLPHQVPAPSHLPQNFLPIAOPAPCSQL
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Bukaryota's Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia: Butheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 1350)
Mu.J., Dillon, D.C., Mitcham, J.L., Harlocker, S.L., Jiang, Y.,
Reed, S.G., Kalos, M.D., Fanger, G.R., Day, C.H., Retter, M.W.,
Stolk, J.A., Skeiky, Y.A., Wang, A. and Mespher, M.J.
Compositions and methods for the therapy and diagnosis of prostate
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/codon_start=1
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1. 1130
Acranism="Homo sapiens",
db_xref="taxon:9606"
/chromosome="19"
/map="19913.3"
42. 356
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                                                                                                                                                                                                                                    /dyanism="Homo sapiens"
/dyaref="taxon:9606"
/dyaref="taxon:9606"
/chromosome="19"
/dyaref="1913"
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/tissue_type="prostate;"
/tissue_type="prostate
/dyas="male"
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in Benign and Malignant Prostate Tissue
1 (Dases 1 to 990)
Moreno, J. (Dases 1 to 990)
Direct Submission
Submitted (08-NOV-1994) Jose M J G Moreno, Urology, Thomas
19107, USA
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Croce.C., Gomella.L., Mulholland,S.Grant., Moreno,J.G. and
Nucleic acid primers for detecting micrometastasis of prostate
cancer
Patent: US 5674682-A 13 07-0cr-1997;
Location/Qualifiers
1. 992
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/product="prostate specific antigen"
93. .113
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291 c 273 g 217 t
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167863.1 GI:2829985
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291 c 272 g
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/protein_id="AAA59995.1"
/db_xref="G1:189524"
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/note="AA 18. .24 propeptide; AA 69 glycosylated Asn
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Homo sapiens prostate-specific antigen mRNA, complete cds.
M26663
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Ref [1] reports bases 1. .135 only.
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361 c 410 g 316 t
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Homo sapiens prostate gland cDNA to mRNA.
Homo sapiens
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/organism="Homo sapiens"
/db_xref="taxon:9606"
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/map="19q13.3"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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1 (bases 1 to 1415)
Riegman, P.H., Klaassen, P., van der Korput, J.A., Romijn, J.C.
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Biochem. Biophys. Res. Commun. 155 (1), 181-188 (1988)
88326297
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CORIXA CORPORATION (US)
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CORIXA CORPORATION (US)
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Unclassified.
1 (bases 1 to 1462)
Croce,C., Gomella,L., Mulholland,S.Grant., Moreno,J.G. and Fischer,R.
Nucleic acid primers for detecting micrometastasis of prostate
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Best Local Similarity 100.0%; Pred. No. 0.012;
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/note="G00-119-695"
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S Nakagawara, H.
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AI926979 wo68d11.x AA654527 nt59a12.s

BF856615 MR1-FT020 BF856642 MR1-FT020

AA420570 nc61b06.r AA228836 nc16h06.r AA226359 nc17h09.r

AA228953 nc14a04.r AA228822 nc15a04.s AA635287 nt71b07.s BF476990 naa58all.

BE838775 RC5-FT019 BE838775 RC6-FN011 BE771925 CM3-FT009

AA654296 nt83c06.s

Perfect score:

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AA79159 nf28b08.8
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AA25115 nc21q11.8
AA574023 nf44f11.8
AA574023 nf44f11.8
AA534235 nf70e05.8
BR856617 MR1-FT020
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similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN)....
mRNA sequence.
AA657663
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Manall: egapbs refemall.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.,
Michael Emmert-Buck, M.D., Ph.D.
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.

(Dases 1 to 189)

NCI-GAP http://www.ncbi.nlm.nlh.gov/ncicgap.

NATional Cancer Institute, Cancer Genome Anatomy Project (CGAP),
Unpublished (1997)
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Bukaryota; Metzoos; Chordata; Craniata; Vertebrata; Buteleostomi;

Mammalia; Eutheria; Primates; Catarrhin; Hominidae; Homo.

Massas 1 to 196)

Massas 2 to 196)

Massas 2 to 196

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BF858362.1 GI:12246106
EST.
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Tel: +55-11-2704922
Tel: +55-11-27001
Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gephtml2.pl?tl=RC5st2=RC5-FT0193-21100-012-G09_1&t3-200-11-21&t4-1)
Seq primer: puc 18 forward
High quality sequence stop: 190.
Location/Qualifiers
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Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
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Troc. Natl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000)
2020263
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100.0%; Score 27; DB 9; Length 189;
Best Local Similarity 100.0%; Pred. No. 0.16;
Matches 27; Conservative 0; Mismatches 0; Indels
Trace considered overall poor quality
Seq primer: -folm15 fwd. ET from Amersham
High quality sequence stop: 1.
Location/Qualifiers
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Prazi +55-11-2704922
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Email: asimpson@lidwig.org.br
Email: asimpson@lidwig.org.br
Email: asimpson@lidwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripte/gethtml2.pl?tl=RC5st2=RC5-FT0193-
211100-012-670543-2000-11-21644=1)
Seq primer: puc 18 forward
High quality sequence stop: 239.
Location/Qualifiers

1. 239
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//dev_stage="Adulit"
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Site_2: SmaI; A mini-library was made by cloning products
derived from ORESTES PR(U.S. Letters Patent application
No. 196,716 - Ludwig Institute for Cancer Research)
profiles into the puc 18 vector. Reverse transcription of
tissue mRNA and cDNA amplification were performed under
1. .196
/organism="Homo sapiens"
/db_xrof="taxon:9606"
/clone_lib="FT0193"
/dev_stage="Adult"
/note="Crgan: prostate_tumor; Vector: pucl8; Site_1: SmaI;
Site_2: SmaI; A mini-library was:made by cloning products
derived from ORESTES PCR (U.S. Letters Patent application
No. 196.716 - Ludwig Institute for Cancer Research)
profiles into the pUC 18 vector. Reverse transcription of
tissue mRNA and cDNA amplification were performed under
low stringency conditions."
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Bukaryota; Metzoo; Chordata; Craniata; Vertebrata; Euteleostomi;

Mamalia; Eutheria; Primates; Catarrhin; Hominidae; Homo.

1 (bases 1 to 239)

1 (bases 1 to 239)

1 (bases 2 to 239)

1 (bases 3 to 239)

1 (bases 3 to 239)

1 (bases 4 to 239)

1 (bases 5 to 239)

1 (bases 6 to 239)

1 (bases 1 to 239)

2 (bases 2 to 239)

2 (bases 2 to 39)

2 (bases 2 to 39)

3 (bases 2 to 39)

4 (bases 2 to 39)

5 (bases 2 to 39)
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FRC5-FF0193-211100-012-G07 FT0193 Homo sapiens CDNA, mRNA sequence.
BFB58359
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Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
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Proc. Natl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000)
20202663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 27; DB 12; Length 196; 100.0%; Pred. No. 0.16;
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ORIGIN

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T29518 272 bp mRNA linear EST 06-SEP-1995 EST828353 Human Prostate gland Homo sapiens cDNA 5' end similar to antigen, prostate specific (HT:2350), mRNA sequence.
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932 Clopper Rd, Gaithersburg, MD 20878
Tel: 3018699056
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fax: 3018699423
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FRAX: #23-11-2./U.V.U.

FRAX: #23-11-2./U.V.U.

This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL

(http://www.ludwig.org.br/scripts/gethtml2.pl?tl-&t2-PM1-FT0028-030
700-001-c02_1&t3=2000-07-03&t4-1)

Seq primer: puc 18 forward

High quality sequence stop: 241.

Location/Qualifiers

1. .24

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/note="Organ: prostate_tumor; Vector: pucl8; Site_1: Smal; Site_2: Smal; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research)

profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
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1 (bass 1 to 241)
Dias Neto, E., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R., Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F., Goldman, G.H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H., Brunstein, A., deoliveira, P.S., Bucher, P., Jongeneel, C.V., O'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and
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Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research:
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shotgun sequencing of the human transcriptome with ORF expressed
                                                                                                          Gaps
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                                                               Length 239;
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                                                           ch 100.0%; Score 27; DB 12; Similarity 100.0%; Pred. No. 0.17; 27; Conservative 0; Mismatches 0;
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+55-11-2707001
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      67
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BE769440
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TITLE

FEATURES

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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Bukaryota; Metazoa; Catarrhini; Hominidae; Homo.

I (bases 1 to 272)

Adams, M.D., Kerlavage, A.R., Fleischmann, R.D., Fuldner, R.A., Bult

C.J., Lee, M., Kirkness, E.F., Weinstock, K.G., Gocayne, J.D., White

O., Sutton, G., Blake, J.A., Brandon, R.C., Chiu, M.-W., Clayton, R.A.,

Cline, R.T., Cotton, M.D., Earle-Hughes, J., Fine, L.D., FtrzGerald

L.M., FitzHugh, M.M., Fritchman, J.L., Geoghagen, N.S.M., Glodek, A.,

Gnehm, C.L., Hannan, M.C., Hedblom, E., HinkleJr, P.S., Kelley, J.M.,

Klimek, K.M., Kelley, J.C., Liut, L.-T., Marmaros, S.M., Merrick, J.M.,

Moreno-Palanques, R.F., McDonald, L.A., Nguyen, D.T., Pellegrino, S.M.,

Bhillips, C.A., Ryder, S.E., Scott, J.L., Saudek, D.M., Shritey, R.,

Small, K.V., Spriggs, T.A., Utterback, T.R., Weidman, J.F., Li, Y.,

Bednarik, D.P., Cao, L., Cepeda, M.A., Coleman, T.A., Collins, E.-J.,

Bumke, D., Feng, P., Ferrie, A., Fischer, C., Hastings, G.A., Hew, W.H.,

Kunsch, C., Ji, H., Li, H., Melssner, P.S., Olsen, H., Kaymond, L., Wei,

Y.-F., Wing, J., Xu, C., Yu, G.-L., Ruben, S.M., Dillon, P.J., Fannon

Worter, T. C., Reader, C.M., and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BE588848 1 CL_CGAP_Br16 Homo sapiens cDNA clone IMAGE:3334468 3/
Similar to SW:PROS_HUMAN P07288 PROSTATE SPECIFIC ANTIGEN PRECURSOR // mRNA sequence.
BF588848 1 GI:11681172
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Oro clone availability, additional sequence and expression
information related to this EST, please contact the TIGR Database
(tdbinfo@tdb.tigr.org)
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/note="Organ: prostate gland"
81 c 71 g 56 t
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/organism="Homo sapiens"
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/db_xref="taxon:9606"
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BF856613

110 MR1-FT0200-311000-001-b11 FT0200 Homeo Sapiens CDNA, mRNA sequence.

S BF856613.1 G1:12244370

S EST

Houna.

ISM Homeo sapiens

ELKALYOLE, Metazca; Chordata; Craniata; Vertebrata; Euteleostomi; Callaryole, Metazca; Chordata; Catarrhin; Hominidae, Homeo

Bransla; Butheria; Primates; Catarrhin; Hominidae, Homeo

S Dias Nato, E., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R., Goldman, G. H., Carvalho, A.F., Mateukuma, A., Baia, G.S., Simpson, D. H., M. J., Soares, P., Brentani, R.R., Reis, L.F., de Souza, S.J., and Shotgun sequencing of the human transcriptome with ORF expressed sequence tags
                                                                                                                                                                                                                              /organiam="Homo sapiens"
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/clone="IMAGE:982382"
/clone=lib="Mori_CGAP_Priz"
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/lasue_type="metastatic prostate bone lesion"
/lab_bost="bhi0B"
/note="vector: pAMP10; mRNA made from metastatic prostate lesion of the bone, CDNA mede by oligo-dr priming.
Non-directionally cloned. Size-selected on agarose gel, average insert size 600 bp. Library made by D. Krizman, NIH-"
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Teal: +55-11-2704922

Email: asimpson@ludwig.org.br

This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project: This entry can be seen in the following URL

(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=MR1&t2=MR1-FT0200-
Seq primer: puc 18 forward

High quality sequence stop: 269.
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Best Local Similarity 100.0%; Score 27; DB 9; Length 310;
Matches 27; Conservative 0; Mismatches 0; Indels 0; Gaps
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Proc. Matl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000)
20202663
Loboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-Sp,
                                                  found through the I.M.A.G.E. Consortium/LLNL at: www.bio.llnl.gov/bbry/image/image.html
Insert Length: 426 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham.
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                                           Homo sapiens

Bukaryota; Metacas; Chordata; Craniata; Vertebrata; Euteleostomi;

Bukaryota; Metacas; Chordata; Craniata; Vertebrata; Euteleostomi;

CERP 1 (bases 1 to 288)

NCI-GGAP http://www.ncbi.nlm.nih.gov/ncicgap.

Tumor Gene Index

Tususberg, Ph.D.

Email: cgapbs.remail.nih.gov/ncicgap.

Contact: Robert Strausberg, Ph.D.

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Tususe Procurement: Dennis Sgroi, M.D., Kristina Cole, M.D., Ph.D.

Contact: Robert Strausberg, Ph.D.

Email: cgapbs.remail.nih.gov

Tususe Procurement: Dennis Sgroi, M.D., Kristina Cole, M.D., Ph.D.

Contact: Robert Strausberg, Ph.D.

Email: cgapbs.remail.nih.gov

Clone distribution: NCI-CGAP clone distribution information can be infoetmage.lin.gov

Clone distribution: NCI-CGAP clone distribution information can be infoetmage.lin.gov

Location/Qualifiers

Location/Qualifiers

Location/Qualifiers

Alab.host="Manage" Homos aspiens"

Alab.host="Manage" Homos aspiens"

Alab.host="Manage" Homos adult"

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Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Rosa, Metazoa; Lo 310)

Nolic Gab Interpi/www.ncbi.nlm.nih.gov/ncicgap.

Nolic Gab Index

Tumor Gene Index

Tumor Gene Index

Tumor Gene Index

Nolic Gap Index

Tumor Gene Index

Talsue Contact: Robert Strausberg, Ph.D.

Email: cgapba-remail.nih.gov

Talsue Procurement: W. Douglas Figg, Ph.D., Paul H. Duray, M.D., CDNA Library Preparation: David B. Krizman, Ph.D.

CDNA Library Arrayad by: Greg Leenon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
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27; Conservative 0; Mismatches 0; Indels (
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AUTHORS
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases I to 329)
Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Brunstein,A., deoliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fai: +55-11-2704922
Fax: +55-11-2707001
Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=RC5&t2=RC5-FT0193-
201100-012-H06&t3=2000-11-20&t4=1)
Seq primer: puc 18 forward
High quality sequence strar: 48
High quality sequence stop: 329.
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                                                                                                                                                                                                                     BF858271 329 bp mRNA linear EST 16-JAN-200
RC5-FT0193-201100-012-H06 FT0193 Homo sapiens cDNA, mRNA sequence.
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Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shotgun sequencing of the human transcriptome with ORF expressed
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100.0%; Pred. No. 0.19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert Strausberg, Ph.D.

Email: cgapbs-remail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.

Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.

Michael Emmert-Buck, M.D., Ph.D.

CDNA Library Preparation: David B. Krizman, Ph.D.

CDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
found though the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html
Seq primer: -40ml3 fwd. ET from Amersham

High quality sequence stop: 298.

I. 328
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AA654296 GI:2590450
Site_2: Smal; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under
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NCI-GGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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/lab_host="DH10B"
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Page 6	Brunstein.A., deoliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and Shotgun sequence tags  JOURNAL Proc. Natl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000)  COMMENT COLECT: Simpson A.J.G.  Laboratory of Cancer Genetics  Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,  gel: +55-11-2704992	Fax: +55-11-2707001 Email: asimpsoneludwig.org.br This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL (Attp://www.ludwig.org.br/scripts/gethtmi2.pl?tl=&t2=CM3-FT0097-300 Seq primer: puc. 18 forward High quality sequence start: 25 High quality sequence stop: 355 FEATURES Location/Oualifiers source Location/Oualifiers	/Organism="Homo sapiens" //Obxef="taxon:9666" //Clone_lib="F10097" //Gev_stage="Adult"	Query Match Best Local Similarity 100.0%; Score 27; DB 12; Length 354; Matches 27; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  Qy 1 AAACTTCAGTGTGAGACCTCCATGTT 27  Db 65 AAACTTCAGTGTGGGACCTCCATGTT 91	AA228953 LOCUS AA228953 AA228953 DEFINITION nC14a04.cl NCLCGAP_Pr1 Homo sapiens cDNA clone IMAGE:1008078 annua sequence. ACCESSION AA228953 VERSION AA228953.1 GI:1851772	VISM VISM ICE ORS		Not-CGAP clone distribution information can be
ACCESSION BEB88775 VERSION BE099775	W H S	JOURNAL Proc. Natl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000)  MEDLINE 20202661.  CONTROLT  CONTROLT  Laboratory of Cancer Genetics  Ludwig Institute for Cancer Research  Brazil  Fai: +55-11-270492  Fai: +55-11-270492  Fai: ssimpson@ludwig.org.br	Project This entry can be seen in the FAPESP/LICR Human Cancer Genome (http://www.ludwig.org.br/scripts/gethtmi2.pl?tl=st2=RC6-FN0114-090 High quality sequence start: 8  FEATURES  Source /organism="Home sapiens" / Ab xref="taxon:9606" / Ab xref="taxon:	/Accellange // Accellange // A	atch cal Similari 27; Cons AAACTTCAGTG	RESULT 12 PET71925 LOCUS LOCUS BE771925 BE771925 BE771925 BE771925 VERSION BE771925 VERSION BE771925 SOURCE Numan. ORGANISM Homo.	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; REFERENCE Mammalia; Eutherla; Primates; Catarrhini; Hominidae; Homo. AUTHORS Dias Neto.E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Soldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,	

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Email: cgapbs-remail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.,
Michael Emmert-Buck, M.D., Ph.D.
CDNA Library Preparation: David B. Krizman, Ph.D.
CDNA Library Preparation: David B. Krizman, Ph.D.
CDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D.
DNA Sequencing by: Mashington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 51 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 1.
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strand cDNA was primed with oligo(dT)17 on 50 ng of
DNAse-treated, total cellular RNA obtained from 5,000-10
/OO microdissected cells histologically determined to be
fully malignant prostate cancer cells. Double-stranded
cDNA was ligated to EcoRI adaptors, 5 cycles of FCR
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Unpublished (1997)
On Nov 12, 1997 this sequence version replaced gi:2559129.
Contact: Robert Strausberg, Ph.D.
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Tumor Gene Index
Dupublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-rémail.nin.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.,
Michael Emmert-Buck, M.D., Ph.D.
CDNA Library Preparation: David B. Krizman, Ph.D.
CDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llni.gov/Dbry/Anage/Anage.html
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strand cDNA was primed with oligo(dT)17 on 50 ng of
DNAse-treated, total cellular RNA obtained from 5,000-10
,000 microdissected, histologically normal prostate
epitchelial cells. Double-stranded cDNA was ligated to
EcoRI adaptors, 5 cycles of PCR applied to the cDNA with
an adaptor-specific primer, and the resulting PCR product
subcloned into pAMP10 by the UDG-cloning method (Life
Technologies). Average insert size is 600 bp. NOTE: Not
directionally cloned. This library was constructed by
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ANTIGEN PRECURSOR (HUMAN);,
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the I.M.A.G.E. Consortium/LLNL at:
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                      www-bio.llnl.gov/bbrp/image/image.html
Seq primer: -28mil revl ET from Amersham
High quality sequence stop: 342.
Location/Qualifiers
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applied to the cDNA with an adaptor-specific primer, and the resulting PCR product subcloned into pAMP10 by the UDG-cloning method (Life Technologies). Average insert size is 600 bp. NOTE: Not directionally cloned. This library was constructed by David Krizman."
                                                                                                                                                            BASE COUNT
ORIGIN
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0; Gaps Query Match
Best Local Similarity 100.0%; Score 27; DB 9; Length 398;
Best Local Similarity 100.0%; Pred. No. 0.2;
Matches 27; Conservative 0; Mismatches 0; Indels ò

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Search completed: November 19, 2002, 05:17:33
Job time : 1545.46 secs

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Sequence 616, App
Sequence 616, App
Sequence 616, App
Sequence 980, App
Sequence 977, App
Sequence 97, App
Sequence 5, Appli
Sequence 1, Appli
Sequence 2, Appli
Sequence 280, App
Sequence 280, App
Sequence 34, Appli
Sequence 34, Appli
Sequence 34, Appli
Sequence 34, Appli
Sequence 603, App
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Sequence 153, App
Sequence 334, App
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111.898 Million cell updates/sec
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2. /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
3. /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
3. /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
3. /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
3. /cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
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4. /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
3. /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
4. /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
3. /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
4. /cgn2_6/ptodata/1/pubpna/US09_PUBCOMB.seq:*
4. /cgn2_6/ptodata/1/pubpna/US00_PUBCOMB.seq:*
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4. /cgn2_6/ptodata/1/pubpna/US00_PUBCOMB.seq:*
GenCore version 5.1.3
Copyright (c) 1993 - 2002 Compugen Ltd.
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US-09-729-143-616
US-09-822-827-616
US-09-822-827-977
US-09-822-827-980
US-09-755-100-3
US-09-755-100-3
US-09-755-100-3
US-09-755-100-3
US-09-755-100-3
US-09-755-100-1
US-09-969-708-280
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US-09-770-445-844
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US-09-759-143-334
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US-09-919-580-34
                                                                                                                                                                                                                                                                                                                                                                   1 aaacttcagtgtgtggacctccatgtt 27
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                      - nucleic search, using sw model
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Maximum DB seq length: 200000000
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Sequence 9, Application 15/09829004A

Sequence 9, Application 15/09829004A

Patent No. US20020132976A1

Sequence 9, Application 15/09829004A

Sequence 9, Application 15/09829

Sequence 100 13/09824

Sequence 100 14.24

TITLE OF INVENTION: Immunogenic Peptides Derived from Prostate-Specific Antigen (P TITLE OF INVENTION: USes Thereof

FILLE OF INVENTION: USes Thereof

FILLE OF INVENTION UNMBER: US/09/829,004A

CURRENT APPLICATION NUMBER: US 60/195,456

PRIOR APPLICATION NUMBER: US 60/195,456

PRIOR FILING DATE: 2000-04-10

NUMBER OF SEQ ID NOS: 12

SEQ ID NO 9

SEQ ID NO 9
                                                                                                                                                                                                                                     6, Appli
3, Appli
1, Appli
1938, Ap
2142, Ap
2083, Ap
    Sequence 334, App
Sequence 638, App
Sequence 698, App
Sequence 698, App
Sequence 37, Appli
Sequence 378, App
Sequence 1379, App
Sequence 245, App
Sequence 245, App
Sequence 245, App
Sequence 245, App
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Sequence 101, App
Sequence 2307, Ap
Sequence 6, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 1938, Ap
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                                                                                                                                              Sequence 321, App
Sequence 322, App
Sequence 899, App
Sequence 12568, A
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899, App
12568, A
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0 US-09-780-669-334

0 US-09-822-827-334

0 US-09-759-143-698

0 US-09-734-698

0 US-09-954-511-985

US-09-954-511-985

US-09-954-511-1379

US-09-960-352-2463

US-09-960-352-13191

US-09-960-352-331

US-09-960-352-331

US-09-960-301-301

US-09-960-301-301

US-09-960-301-301

US-09-960-301-301

US-09-960-301-301

US-09-960-301-301

US-09-976-4801-1338

US-10-041-395-4

US-10-041-395-4

US-09-976-869-2083
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ch 100.0%; Score 27; DB 10; Similarity 100.0%; Pred. No. 0.00059; 27; Conservative 0; Mismatches 0;
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Patent No. US200202248A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
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Dillon, Davin C.
Mitcham, Jennifer
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US-09-759-143-616
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APPLICANT:

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US-09-822-827-616
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APPLICANT: Stolk, John A.
APPLICANT: Day, Craig H.
APPLICANT: Day, Craig H.
APPLICANT: Carter, Darrick
APPLICANT: Carter, Darrick
APPLICANT: Carter, Darrick
APPLICANT: Mang, Aljun
APPLICANT: Mang, Aljun
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Skeiky, 1111am
TITLE OF INVENTION: COMPOSITIONS OF PROSTATE CANCER
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
TITLE OF INVENTION UNMER: US/09/759,143
CURRENT FILING DATE: 2001-01-12
NUMBER OF SEQ ID NOS: 934
SOFTWARE: FASISEQ for Windows Version 3.0
SEQ ID NO 616
LEBNOTH: 1350
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Hival, John
APPLICANT: Hival, John
APPLICANT: McNeill, Patricia D.
APPLICANT: Houghton, Raymond L.
APPLICANT: Houghton, Raymond L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
CURRENT APPLICATION NUMBER: US/09/780,669
CURRENT APPLICATION NUMBER: 2001-02-09
WUMBER OF SEQ ID NOS: 943
SOFFWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ouery Match

100.0%; Score 27; DB 10;
Best Local Similarity 100.0%; Pred. No. 0.0013;
Matches 27; Conservative 0; Mismatches; 0;
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; Patent No. US20020051977A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Xu, Jiangchun
APPLICANT: Mitcham, Jennifer L.
APPLICANT: Hitcham, Jennifer L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Harlocker, Susan L.
APPLICANT: Henderson Robert A.
APPLICANT: Retter, Marc W.
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APPLICANT: Roll, John A.
APPLICANT: Veduvick, Thomas S.
APPLICANT: Carter, Marc W.
APPLICANT: Carter, Marc W.
APPLICANT: Gary R.
APPLICANT: Li, Samuel
Harlocker, Susan L.
Jiang, Yuqui
Henderson, Robert A.
Ralos, Michael D.
Fanger, Gary R.
Retter, Marc W.
Stolk, John A.
Day, Craig H.
Vedvick, Thomas S.
Carter, Thomas S.
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Skeiky, Yasir A.W.
Hepler, William
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; ORGANISM: Homo sapien
US-09-759-143-616
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; ORGANISM: Homo sapien
US-09-780-669-616
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| Beat Local Stanlarity 100.04; Score 27; DB 10; Length 1350; Beat Local Stanlarity 100.04; Score 27: DB 10; Length 1350; Beat Local Stanlarity 100.04; Score 20: DB 1120 AACTTCAGGGGCCTCCANGTT 1155
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| BESULT 4 | BAACTTCAGGGCCTCCANGTT 1155
| BESULT 4 | BAACTTCAGGGCCTCCANGTT 1155
| Sequence 6.16, Apilcation US/09822827 | BESULT 4 | BAACTTCAGGGCTTCANGTT 1155
| Sequence 6.16, Apilcation US/09822827 | BACTTCAGGGCTTCANGTT 1155
| Sequence 6.16, Apilcation US/09822827 | BACTTCAGGGCTTCANGTT 1155
| Sequence 6.16, Apilcation US/09822827 | BACTTCAGGGCTTCANGT 1155
| STATE CO TANGTTON UNMARKED 100 UN SCORE 27; DB 10; Length 1350; BACTTCAGGGCTTCANGT 1155
| STATE CO TANGTTON UNMARKED 100 UN SCORE 27; DB 10; Length 1350; BACTTCAGGGCTTCANGT 1155
| SCOPTHARE: Pack SEC 100 UN SCORE 27; DB 10; Length 1350; BACTTCAGGGCTTCANGT 1155
| BACTTCAGGGCTTCANGGCTTCANGGCTTCANGT 1155
| BACTTCAGGTGTGGACCTCCANGT 1155
| BACTTCAGGGTGTGGACCTCCANGT 1155
| BACTTCAGGGTGTGACCTCCANGT 1155
| BACTTCAGGGTGTGGACCTCCANGT 11155
| BACTTCAGGGTGTGACCTCCANGT 1155
| BACTTCAGGGTGTGACCTCCANGT 1155
| BACTTCAGGGTGT
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Gaps

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Sequence 3, Application US/09755100
Fatent No. US20020099189A1
GENERAL INFORMATION:
APPLICANT SAVITAKY, Kinneret et al.
TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REFERENCE: 2786-0156P
CURRENT APPLICATION NUMBER: US/09/755,100
CURRENT FILING DATE: 1999-01-01-08
PRIOR APPLICATION NUMBER: IL 128587
PRIOR APPLICATION NUMBER: IL 129439
PRIOR FILING DATE: 1999-04-14
PRIOR PILING DATE: 1999-04-14
PRIOR PLILING DATE: 1999-04-14
PRIOR PLILING DATE: 1999-04-14
PRIOR PLILING DATE: 1999-08-11
SUMBER OF SEQ ID NOS: 12
SOFTWARE: PATENTIN VEY: 2.1
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Patent No. US2002009918911

GENERAL INFORMATION:

APPLICANT: SAVIT&X, Kinneret et al.

TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES

FILE REPERENCE: 2786-0156P

CURRENT APPLICATION NUMBER: UL 128587

PRIOR APPLICATION NUMBER: UL 129439

PRIOR PELING DATE: 1999-04-14

PRIOR FILING DATE: 1999-04-14

SEQ ID NOS: 12

SOFTWARE: PATENTIN VOICE. 2.1

SEQ ID NO 1

LENGTH: 4661
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                                                                                                                                                   100.0%; Score 27; DB 10; Length 3423; 100.0%; Pred. No. 0.0016;
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100.0%; Score 27; DB 10;
Best Local Similarity 100.0%; Pred. No. 0.0017;
Matches 27; Conservative 0; Mismatches 0;
                                                                                                                                                                                                         0; Mismatches
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                                                                                                                                                         Query Match 100.(
Best Local Similarity 100.(
Matches 27; Conservative
                                                   TYPE: DNA
CORGANISM: Homo sapiens
US-09-755-100-5
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US-09-755-100-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 3
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US-09-755-100-3
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     SEQ ID NO 5
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Sequence 980, Application US/09822827

Patent No. US20020081680A1

GENERAL INFORMATION:

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER

FILE REFERENCE: 210121.534C1

CURRENT APPLICATION NUMBER: US/09/822,827

WUMBER OF SEQ ID NOS: 982

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 980

LENGTH: 2406
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 5, Application US/09755100
Patent No. US20020099189A1
GENERAL INFORMATION:
TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REFERENCE: 2786-0156P
CURRENT APPLICATION NUMBER: US/09/755,100
PRIOR APPLICATION NUMBER: US/09/755,100
PRIOR PAPLICATION NUMBER: IL 128587
PRIOR PAPLICATION NUMBER: IL 129439
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-04-14
PRIOR FILING DATE: 1999-04-14
PRIOR FILING DATE: 1999-04-14
                  Sequence 977, Application US/09822827
Batent No. US20020081680A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.534C1
CURRENT APPLICATION NUMBER: US/09/822,827
CURRENT FILING DATE: 2001-03-28
NUMBER OF SEQ ID NOS: 982
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 977
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100.0%; Score 27; DB 10;
Best Local Similarity 100.0%; Pred. No. 0.0015;
Matches 27; Conservative 0; Mismatches 0;
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SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                          ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-822-827-977
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-822-827-980
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US-09-822-827-977
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LOCATION: 389, 921, 122, 123, 295, 320, 322, 352, 355, 361, 369, 375, 10.0CATION: 389, 921, 122, 123, 295, 320, 322, 352, 355, 361, 440, 445, 10.0CATION: 382, 384, 386, 389, 409, 411, 413, 414, 415, 421, 440, 445, 10.0CATION: 448, 453, 461, 500, 510, 528, 532, 537, 546, 547, 563, 564, 10.0CATION: 569, 577, 583, 587, 597, 598, 599, 609, 613, 616, 621

COTHER INFORMATION: n = A,T,C or G

NAME/KEY: misc_feature

LOCATION: 622, 627, 648, 681

COTHER INFORMATION: n = A,T,C or G

US-09-919-580-34
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Patent No. US20020110832a1

GENERAL INFORMATION:
APPLICANT: Pyle, Ruth
APPLICANT: Xu, Jiangchurh
APPLICANT: Xu, Jiangchurh
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 20121.552

CURRENT APPLICATION NUMBER: US/09/919,580.
CURRENT FILING DATE: 2001-07-30

NUMBER OF SEQ ID NOS: 934

SEQ ID NO 34

SEQ ID NO 34

LENGTH: 684
                                                                                                          Sequence 72, Application US/09919580
Fatent No. US20020110832A1
GENERAL INFORMATION:
APPLICANT: FULL
APPLICANT: With
APPLICANT: Such 31 angehun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLÓN CANCER
FILE REFERENCE: 2011.1.552
CURRENT FILING DATE: 2001-07-30
NUMBER OF SEQ ID NOS: 934
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 72.6%; Score 19.6; DB 10; Length 684; Best Local Similarity 84.6%; Pred. No. 3.6; Matches 22; Conservative 0; Mismatches 4; Indels 0;
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72.6%; Score 19.6; DB 10; Length 601;
Best Local Similarity 84.6%; Pred. No. 3.5;
Matches 22; Conservative 0; Mismatches 4; Indels 0
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ORGANISM: Homo sapiens
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-919-580-72
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US-09-919-580-34/c
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US-09-969-708-280/C
US-09-969-708-280/C
Sequence 280, Application US/09969708
Fetent No. US20020102532A1
GENERAL INFORMATION:
APPLICANT: Augustus, Meena
TITLE OF INVENTION: Setens
TITLE OF INVENTION: Setens
FILE REFERENCE: 689290-70
CURRENT FILING DATE: 2000-10-03
FRICH REPLICATION NUMBER: US/60/237,606
FRIOR PILING DATE: 2000-10-03
FRIOR APPLICATION NUMBER: US/60/237,608
FRIOR FILING DATE: 2000-10-03
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US-09-755-100-2

Sequence 2, Application US/09755100

Patent No. US20020099189A1

GENERAL INFORMATION:

TITLE NEVERALISMENCE: 2786-0156

TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES

TITLE OF INVENTION: NOVER: US/09/755,100

CURRENT APPLICATION NUMBER: US/09/755,100

CURRENT APPLICATION NUMBER: IL 128587

PRIOR FILING DATE: 1999-02-18

PRIOR FILING DATE: 1999-04-14

NUMBER OF SEQ ID NOS: 12

SEQ ID NO 2

LENGTH: 4661

LENGTH: 4661

LENGTH: 4661

STORMANES: Homo sapiens

US-09-755-100-2
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                                      Indels
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Pred. No. 0.0017;
; Mismatches 0;
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                                                                                               178 AACTTCATTGTTTGGACCTGCATATT 153
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       Best Local Similarity 100.0%; Pi
Matches 27; Conservative 0;
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US-09-969-708-280
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72.6%; Score 19.6; DB 10; Length 1168;
Best Local Similarity 84.6%; Pred. No. 4;
Matches 22; Conservative 0; Mismatches 4; Indels 0;
US-09-925-300-603

Sequence 603, Application US/09925300

Patent No. US20020151681A1

GENERAL INFORMATION:

APPLICANT: Card Rosen,

APPLICANT: Steve Ruben

TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies

FILE REFERENCE: PAIO.

CURRENT APPLICATION NUMBER: US/09/925,300

CURRENT FILING DATE: 2001-08-10

PRIOR APPLICATION NUMBER: PCT/US00/05988

PRIOR PILING DATE: 1000-03-08

PRIOR APPLICATION NUMBER: 60/124,270

PRIOR PILING DATE: 1999-03-12

NUMBER OF SEQ ID NOS: 1890

SOFTWARE: PatentIN Ver. 2.0

SEQ ID NO 603

LENGTH: 1168

TYPE: DNA

ORANISM: Homo sapiens

FEATURE:

NAME/KET misc feature

LCCATION: (1121)

OTHER INFORMATION: n equals a,t,g, or c

LCATION: (1123)

OTHER INFORMATION: n equals a,t,g, or c

LCATION: (1133)

OTHER INFORMATION: n equals a,t,g, or c

LCATION: (1153)

COTHER INFORMATION: n equals a,t,g, or c

LCATION: (1153)

COTHER INFORMATION: n equals a,t,g, or c

LCATION: (1153)

COTHER INFORMATION: n equals a,t,g, or c

LCATION: (1153)

COTHER INFORMATION: n equals a,t,g, or c
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24 1 atgtgggtcccggttgtcttcctc 24 IDENTITY\_NUC Gapop 10.0 , Gapext 1.0 Perfect score: Sequence: Scoring table:

4109280 2054640 seqs, 14551402878 residues Searched:

Total number of hits satisfying chosen parameters:

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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em\_in:\* em\_mu:\* em\_om:\* em\_ov:\* em\_pat: em\_ro:\* em\_or:\*

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em\_htg\_mam:\* em\_htg\_vrt:\* em\_sy:\* em\_htgo\_hum:\* em\_htg\_mus:\* em\_htg\_pln:\* em\_htg\_rod:\*

Pred. No. is the number of results predicted by chance to have a

em\_htgo\_other:\*

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description		717	AKI4/230 Sequence	CARAGE DENIMINERS 9	AT3109	AJ459783 Homo sapi	AJ4597	C17040 Human prost	: 167863 Sequence 13	A.J.45971	, AF3354	, M26663 Homo sapien	167864 Sequence 14	£32812	X05332 Human; mKNA	7. BC005307 HOMO Sapi	COSTOSS COMPONE	AF335478 HOMO sani	AX106218 Sequence	AX 8334	. AX0334	PX0334	AX0334	X14810 Human DNA f	M24543 Human prost	A37262 Sequence 2	AR167395 Sequence	uman F	ACOLIDZS HOMO SAPI	AF243527 Homo sapi				22	AX434375 Sequence		luman D	НОШО	AC115837 Homo sapi	ACUU9629 HOMO	Homo	1111	AJ4094/9 EMELLUELL	
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T VICTOR	ID		AX281250	AK14/236	CJEJEE	HSA310937	HSA459783	HSA459782	HSU17040	167863	HSA459784	AF335477	HUMAPS	I67864	E32812	HSPSAR	BC005307	AY033403	AF335478	AX106218	AX033404	AX033402	AX033400	AX033401		HUMPSANTIG	A37262	AR167395	HUMPSAA		AF243527	MMPROS	7	33	5	AX434375	7	25	561	AC115837	200	K09308	200	7 5 6 0 7	
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ALIGNMENTS

linear PAT 02-NOV-2001

RESULT 1 AX281250	
LOCUS	AX281250 24 bp DNA
DEFINITION	Sequence 7 from Patent WO0176622.
ACCESSION	AX281250
VERSION	AX281250.1 GI:16608506
KEYWORDS	
SOURCE	synthetic construct.
ORGANISM	synthetic construct
	artificial sequences.
REFERENCE	
AUTHORS	Pedyczak, A., Chong, P. and Sia, C.D.
TITLE	Immunogenic peptides derived from prosta

ate-specific antigen (psa) and uses thereof Patent: WO 0176622-A 7 18-OCT-2001;

JOURNAL

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prim_transcript
mRNA
                                                                                              TATA_signal
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Digby, M.R.

Direct Submission

Submitted (09-JAN-1989) Digby M.R., The Howard Florey Institute,
Melbourne University, Parkville, Victoria 3052, Australia

(Dassa 1 to 389)

Digby, M., Zhang, X. Y. and Richards, R.I.

Human prostate specific antigen (PSA) gene: structure and linkage to the kallikrein-like gene, hGK-1

Nucleic Acids Res. 17 (5), 2137 (1989)
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Mammaliai Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 389)
                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                Unknown.
Unclassified.
Unclassified.
1 (bases 1 to 70)
Everhart, D.S., Kaylor, R.M. and McGrath, K.
Everhart, D.S., Kaylor, R.M. and McGrath, K.
Patterned binding of functionalized microspheres for optical
diffraction-based blosensors
Patterned 5621579-A 2 24-APR-2001;
Location/Qualifiers
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                                                                                                                                             Length 24;
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kallikrein; prostate specific antigen; PSA gene.
Homo sapiens.
Homo sapiens.
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Location/Qualifiers
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                              1. 24
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1 a 7 c 7 g 9 t
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Sequence 2 from patent US 6221579.
AR147236 GI:15111039
Aventis Pasteur Limited (CA)
Location/Qualifiers
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569 bp mRNA linear PRI 15-JUN-1995
PSA-prostate-specific antigen [human, breast cancer specimen, mRNA
partial, 569 nt].
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FQVSHSFPEPPLVDWSLLKRNRFLERPGDDSSHDLMILRLSEPAELTDAVKVWDLPTQEPA
IGTTCYASGWGSIEPEEFLIPRKKLQCVDL"
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Manmalia: Eutheria; Primates; Catarrhini; Hominidae; Homo.

Monne, M., Croce, C. M., Yu, H. and Diamandis, E. P.

Molecular characterization of prostate-specific antigen messenger cancer Res. 54 (24), 6344-6347 (1994)

7537255
GenBank staff at the National Library of Medicine created this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       General staff at the National Library of Medicine created this entry [NCBI gibbsq 161505] from the original journal article. Location/Qualifiers
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Best Local Similarity 100.0%; Pred. No. 0.67;
Matches 24; Conservative 0; Mismatches 0; Indels
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273. ->18
273. ->18
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hotte="prostate-specific antigen"
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232. .318
232. .318
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S75755.1 GI:861469
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Eur. J. Blochem. 268 (16), 4408-4413 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                          Direct Submission
Submitted (11-APR-2001) Courty Y., Faculte de Medecine, EMI-U 0010,
laboratoire d'Enzymologie, 2 bis bd Tonnelle, 37032 Tours cedex,
                                                                                                                                                                                      PRI 17-JAN-2002
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                                                                                                                                                                                                                                                                               Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                   Gaps
                                                                                                                                                                                                                                                      alternative splicing; HKLK3 gene; kallikrein 3; preproPSA-RP2
                                                                                                                                                                                    HSA310937 708 bp mRNA linear PRI 1'
Homo sapiens mRNA for putative preproPSA-RP2 (KLK3 gene),
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                                        9; Length 569;
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                                                                 0; Indels
                                      Score 24; DB 9;
Pred. No. 0.67;
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                                                                 0; Mismatches
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/db_xref="taxon:9606"
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/cell_line="LNCaP"
1. 708
/gene="KLK3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               related accession number AJ310938.
Location/Qualifiers
1. 708
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 160
                                                                                                         42 ATGTGGGTCCCGGTTGTCTTCCTC 65
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                                     Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 24; Conservative 0
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38. .580
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Courty, Y.
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21393944
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22. 807

22. 807

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CNGYLQGITSWGSEPCALPERRSLYTKVVHYRKWIKDTIVANF"
                                                                                                                                                                                                                                                                                                                HSA459783 870 bp mRNA linear PRI 09-MAY-2002
Homo sapiens mRNA for prostate specific antigen (KLK3 gene), splice
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (03-MAY-2002) Courty Y., EMI-U 0010, Laboratoire d'Enzymologie, Universite F. Rabelais, 2 bis byd Tonnelle, Tours, 37032 cedex, FRANCE alternative splice variant sequences: AJ459782, AJ459784.

Location/Qualifiers
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ch 100.0%; Score 24; DB 9; Length 708; 1 Similarity 100.0%; Pred. No. 0.67; 24; Conservative 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Pred. No. 0.66;
Matches 24; Conservative 0; Mismatches 0; Indels
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257 c 242 g 186 t
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/clone="YC140405-00"
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94. .804
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Courty, Y.
Direct Submission
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genes KLK2

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42. 356
/note="PSA: KLK3; alternatively spliced"
/note="PSA: KLK3; alternatively spliced"
/note="PSA: KLK3; alternatively spliced"
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/product="AaL73204 1"
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HOMO sapiens prostate-specific antigen mRNA, complete cds.
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prostate-specific antigen.
Homo sapiens prostate gland cDNA to mRNA.
Homo sapiens
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                             Direct Submission
Submitted (11-JAN-2001) Compugen Ltd., 72 Pinchas Rosen, Tel Aviv
69512, Israel
                                                                                                                                                  (bases 1 to 1130)
David, A., Engel, S., Azar, I., Bernstein, J., Rotman, G., Savitsky, K.
and Mintz, L.
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Henttu.P. and Vihko.P.
CDNA coding for the entire human prostate specific antigen shows coding for the entire human tissue kallikrein genes high homologies to the human tissue kallikrein genes Biochem. Biophys. Res. Commun. 160 (2), 903-910 (1989) 28246551 2470373
On Jan 5, 1995 this sequence version replaced gi:341511.
Ref [1] reports bases 1. 135 only.
Unusual alternative splicing within the human kallikrein and KLK3 gives rise to novel prostate-specific proteins J. Biol. Chem. 277 (20), 18084-18090 (2002)
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1. .1446
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1. 1130
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/db_rare="taxon:9606"
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/clone="PSA-20"
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/note="G00-119-695"
25. .810
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                                                                                                                       HSA459784 11095 bp mRNA linear PRI 09-MAY-2002 Homo sapiens mRNA for prostate specific antigen (KLK3 gene), splice
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PSPDRELGSFLSLPAPLQAHTPSPSILQQSSLPHQVPAPSHLPQNFLPIAQPAPCSQL
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Submitted (03-MAY-2002) Courty Y., EMI-U 0010, Laboratoire
Submitted (03-MAY-2002) Courty Y., EMI-U 0010, Laboratoire
Submitted (03-MAY-2002) Courty Y., EMI-U 0010, Laboratoire
37032 cedex, FRANCE
alternative splice variant sequences: AJ459782, AJ459783.
Location/Qualifiers
1. 103
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                                                                                                                                                                                                                                                                                                                       Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AJ459784.1 GI:20520644
AJternative splicing; KLK3 gene; prostate specific antigen.
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/note="alternative splice variant 3"
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AF335477.1 GI:18478569
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Courty, Y.
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            Geguence 14 from patent US 5674682.
167864 1 GI:2829986
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CE Unknown.

CE 1 (bases 1 to 1462)

RS Groce, C., Gomella, L., Mulholland, S. Grant,, Moreno, J. G. and Fischer, R.

Nucleic acid primers for detecting micrometastasis of prostate cancer

NAL Patent: US 567482-A 14 07-0CT-1997;

Location/Qualifiers

Ource 1. 1462

Location/Qualifiers

A organism="unknown" 343 a 374 c 422 g 323 t
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/note="AA 18. .24 propeptide; AA 69 glycosylated Asn
residue"
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Best Local Similarity 100.0%; Score 24; DB 9; Length 1446;
Matches 24; Conservative 0; Mismatches 0; Indels 0; Gaps
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100.0%; Score 24; DB 6; Length 1462;
Best Local Similarity 100.0%; Pred. No. 0.66;
Matches 24; Conservative 0; Mismatches 0; Indels (
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/note="G00-119-695"
1 372 c 419 g 322 t
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Perfect score:
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Listing first 45 summaries
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       AAD21361
AAT61707
AAT35293
ABN36013
AAA63735294
ABL60747
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AAT35294
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Prostate specific Prostatic specific Prostatic specific Forward primer F1 Human spliced tran Nucleic acid used Prostate specific CDNA for prostate Forward primer F2 Human prostate exp
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## ALIGNMENTS

RESULT 1 AAD21361

AAD21361 standard; DNA;

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(AVET ) AVENTIS PASTEUR LTD
                            10-APR-2000; 2000US-195456P
                                                                                                                                                                              Unidentified.
                                                                                                                                                                                           Prostate-specific antigen; PSA; immunogénic peptide; immune response; therapy; cancer; prostate cancer; tumour metatasis; prophylaxis; ds.
                                               10-APR-2001; 2001WO-CA00473
                                                                   18-OCT-2001
                                                                                                                                                                                                                                                    28-JAN-2002 (first entry)
                                                                                       WO200176622-A2
                                                                                                                                                                                                                              Prostate specific antigen (PSA) derived peptide, CLP313 encoding DNA.
                                                                                                                  /product= "PSA derived peptide, CL
/note= "CDS does not include stop
                                                                                                                                                           Location/Qualifiers
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Chong P,

Sia CDY;

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RESULT 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The patent discloses immunogenic peptides of prostate-specific antigen (PSA) and nucleic acids encoding them. The peptides of the invention are used to prepare a medicament to elicit an immune response in an animal. They are used to treat cancer such as prostate cancer and tumour metastasis. They are also useful for prophylaxis, for preparing monoclonal or polyclonal antibodies, and in conventional techniques of immunology, molecular biology, cell biology and recombinant DNA technology. The present sequence is a DNA encoding PSA derived peptide, CLP313.
                     Claim 6; Page 38; 49pp; English.
                                          Treatment of benign prostatic hyperplasia or prostatic cancer using an anti-sense oligo:nucleotide targeted against prostate specific antigen or the probasin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT61707;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT61707 standard; cDNA; 27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New polypeptide, useful for treating comprises prostate-specific antigen
                                                                                                                       Zamecnik PA;
                                                                                                                                                                       20-SEP-1995;
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DB; AAE13023.
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ilarity 100.0%;
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derived peptide -
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cc comprises administering to the patient a therapeutic amount of a comprises administering to the patient a therapeutic amount of a cc composition comprising an antisense oligonucleotide which selectively hybridises to prostate specific antigen (PSA) gene or mRNA or to a c probasin gene or mRNA sequence of the patient, where the antisense coligonucleotide inhibits expression of the sequence. The present c sequence represents a specifically claimed antisense oligonucleotide can be used to effectively inhibit the growth of (and for the killing of) c where the state of the patient of the killing of) c where the sequence of the present of the used to effectively inhibit the growth of (and for the killing of) c hyperplastic cells or cancerous cells or prostate cells of the the properties of the prostation of the PSA and probasin genes is specific to prostate cells so that the cantisense oligonucleotide can be administered systemically, making it particularly useful in late stage prostatic cancer which has metasoned the particularly and in which the cells have become resistant to oestrogen cor anti-androgen therapy. The method can also be used in benign prostate hyperplasia, or early stage prostatic cancer, as an alternative for the core radical procedures currently used, such as transurethral resection, creative the prostate concer andical prostatectomy, or physical or chemical castration.
Sequence 27 BP; 9 A; 7 C; 9 G; 2 T; 0 other;
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Matches Query Match Best Local : AAT35293 standard; DNA; 27 Local l Similarity 24; Conserv 100.0%; ilarity 100.0%; Conservative 0 45 ВP 0; Score 24; Pred. No. Mismatches 0.16; DB 18; 0; Length Indels 27; 0, Gaps

0;

В

11-MAR-1997 AAT35293; baculovirus Forward primer F1 for prostate specific antigen. chain reaction; primer; transfer vector; (first entry) PCR; PSA; ss. amplify; prostate specific

Synthetic.

EP725139-A2

31-JAN-1996; 07-AUG-1996 96EP-0101304.

28-APR-1995; 03-FEB-1995; 95US-0430498. 95US-0383111.

(FARB ) BAYER CORP.

Barnett TR, Ng PC, Yeung KK;

WPI; 1996-356106/36.

Recombinant fragments tests prostate-specific antigen (PSA) and immunological for use as an immunoassay calibrator in commercial PSA

Claim 3; Page 5; 19pp; English.

AAT35293-T35295 represent amplification primers for the prostate specific antigen (PSA). These primers were used to insert the CDNA for PSA into a BamHI-NotI-cleaved pVL1939 baculovirus transfer vector. The vector that was produced is ATCC VR-2496. The rPSA can produced by the vector can be used as an immunoassay calibrator. The immunoassays that use the rPSA are for the quantitative determination of PSA in a biological sample. In the assay, the test sample is combined with a

novel method has been developed for treating a patient diagnosed

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messenger RNAs that populate a (sub-)transcriptome, where the CC (sub-)transcriptome comprises messenger RNAs transcribed from multiple CC transcription units that populate a genome. The library comprises CC several oligonucleotides, each capable of hybridising selectively to a set of messenger RNAs transcribed from a given transcription unit of CC the genome, which encodes one or more messenger RNA splice variants. The oligonucleotide libraries are useful for detecting RNAs from a biological sample, in expression profiling studies, in qualitatively or quantitatively characterising the corresponding transcriptome, and in CC detecting RNA transcripts and splice variants of human or animal CC transcriptomes. The libraries may also be used as specialised mini CC libraries to detect transcripts of a sub-transcriptome under a CC particular biological or pathological state, and so allowing the CC detection of tissue and pathology-specific genes such as those genes conjuction; to detect developmental specific pathological condition; to detect developmental specific pa
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02-MAY-2001; 2001US-287724P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          developmental-specific genes
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligonucleotide library;
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0.16;
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Query Match

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Best Local
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                                 analyte can be detected by analysing the diffraction pattern of the transmitted light. The method of the present invention may used to detect and quantify a variety of analytes, e.g. midropranisms, viruses, nucleic acids, hormones, steroids, antibodies, drugd, metals and antigens, either for clinical diagnosis of infections, cancer-and allergies or for detecting microbial contamination of garments. The present sequence is a synthetic oligonucleotide, which was used as an analyte to illustrate the
                                                                                                                          The present invention relates to a method for detecting and quantifying the presence or amount of an analyte within la medium. The method comprises micro-contact printing of analyte specific receptors onto polymer film. The target analyte binds to the polymer film, and results in light diffraction when light is transmitted through the film. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N.B. The sequence data for this patent did no specification, but was obtained in electronic at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                             infections and cancer, comprises reacting analyte with diffraction-enhancing material carrying. specific receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Optical diffraction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA63192 standard; DNA;
                                                                                                                                                                                                      Example 2; Page 23; 38pp; English.
                                                                                                                                                                                                                                                       Diffraction-based analyte detection, useful for clinical diagnosis
                                                                                                                                                                                                                                                                                                                                  (KIMB ) KIMBERLY-CLARK WORLDWIDE INC.
                                                                                                                                                                                                                                                                                                                                                                                     22-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                              15-JUN-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   diffraction-based
nation; infection;
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                                                                                                                                                                                                                                                                                                          Kaylor
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer;
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Pred. No.
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RESULT 6
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Matches
Query Match
                                        The invention provides a method of determining prostate specific antigen (pPSA) in a sample which involves determining the amount of complex formed on treating a sample with a pPSA specific antibody. The method is useful for detecting prostate cancer and distinguishing it from benign prostatic disease. The present sequence represents a DNA encoding a partial fragment of the PSA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sig_peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Prostate specific antigen; PSA; prostate; cancer; prostatic disease; gene; ds.
                    Sequence 90
                                                                                                                                             Determining prostate specific antibody (pPSA), comprises determining the amount of complex formed on treating a sample with a pPSA specific antibody, useful for detecting prostate cancer and distinguishing it from benign prostatic disease
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                                                                                                                         Disclosure; Fig 1; 30pp; English.
                                                                                                                                                                                                                                         Mikolajczyk SD, Rittenhouse HG, Wang TJ,
                                                                                                                                                                                                                                                              (MIKO/) MIKOLAJCZYK S D.
(RITT/) RITTENHOUSE H G.
(WANG/) WANG T J.
(WOLF/) WOLFERT R L.
                                                                                                                                                                                                                                                                                                                        30-APR-1999;
                                                                                                                                                                                                                                                                                                                                              23-FEB-2001; 2001US-0792534
                                                                                                                                                                                                                                                                                                                                                                    18-APR-2002
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24; Conservative
                      BP; 8 A; 26 C; 31 G; 25 T; 0 other;
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/note= "pa
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 Length 90;
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             AAT35294;
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                                 AAT35294 standard; DNA; 93 BP
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Best Local Similarity Matches 24; Conserv
                                                                                                                                                                                                                                cDNA sequences AAT92974-77 encode the prostate specific antigen (PSA). The detection of PSA mRNA in peripheral blood is associated with metastatic prostate cancer. Detection of PSA mRNA in the bloodstream indicates that prostate cells are circulating in the blood and confirms diagnosis of cancer. Circulating prostate cells also indicate a risk of a secondary tumour or bone metastatic anchoring. Detection of the PSA mRNA after prostatectomy or radiotherapy indicates a risk that the prostate cancer has spread and that the surgery and radiotherapy was not effective. PCR primers AAT92971-73 were used for the monitoring and diagnosis of patients with prostate cancer. The method comprises the detection of PSA mRNA by reverse transcriptase polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Diagnosis; prostate cancer; prostate specific antigen; PSA; metastatic prostate cancer; prostate cell; secondary tumour; bone metastatic anchoring; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Monitoring or diagnosis of prostate cancer - by detecting prostate-specific antigen mRNA using specific primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cDNA for prostate specific antigen (PSA).
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                                                                                                                                                                       Sequence 92 BP; 14 A; 28 C; 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 13; 25pp; English.
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38 ATGTGGGTCCCGGTTGTCTTCCTC 61
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                                                                                   l Similarity
24; Conserv
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                                                                                                          Score 24;
Pred. No.
                                                                                                                                                                          G; 22 T;
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RESULT 9
ABV07911
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                                                                                                                                                                                                                     Query Match
Best Local :
                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                        AAT35293-T35295 represent amplification primers for the prostate specific antigen (PSA). These primers were used to insert the cDNA for PSA into a BamHI-NotI-cleaved pVL1393 baculovirus transfer vector. The vector that was produced is ATCC VR-2496. The rPSA can produced by the vector can be used as an immunoassay calibrator. The immunoassays that use the rPSA are for the quantitative determination of PSA in a biological sample. In the assay, the test sample is combined with a labelled conjugate and the response of the conjugate is compared to a standard curve established by performing the assay on one or more calibrators using rPSA (or an immunologically reactive fragment).
                                                                  Human; prostate pharmacogenomic
                                                                                                                                        ABV07911 standard; cDNA; 171 BP
                                                                                                                                                                                                                                            Sequence 93
                                                                                                                                                                                                                                                                                                                                              Claim
                                                                                                                                                                                                                                                                                                                                                                     Recombinant fragments -
                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-356106/36
                                                                                                                                                                                                                                                                                                                                                                                                          Barnett
                                                                                                                                                                                                                                                                                                                                                                                                                                        28-APR-1995;
03-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EP725139-A2
        20-FEB-2001; 2001WO-US05171.
                       23-AUG-2001.
                                      WO200160860-A2
                                                                                                           13-SEP-2002
                                                                                                                           ABV07911;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             baculovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  11-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                         (FARB )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Polymerase chain
                                                                                                                                                                                16
                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                              ب
                                                                                                                                                                                             ATGTGGGTCCCGGTTGTCTTCCTC 24
                                                                                                                                                                                                                                                                                                                                             3; Page 5; 19pp; English.
                                                                                            prostate
                                                                                                                                                                                                                                                                                                                                                                                                                         BAYER CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer F2
                                                                                                                                                                                                                                           BP; 12 A; 23 C;
                                                                                                                                                                                                                                                                                                                                                                   prostate-specific antigen for use as an immunoassay
                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                         Ng
                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                    marker;
                                                                                           expression
                                                                                                                                                                                                                                                                                                                                                                                                                                        95US-0430498
95US-0383111
                                                                                                                                                                                                                                                                                                                                                                                                                                                               96EP-0101304
                                                                                                                                                                                                                                                                                                                                                                                                         PC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for prostate specific antigen.
                                                                                                                                                                                                                     100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             vector;
                                                                                                                                                                                                                                                                                                                                                                                                         Yeung
                                                                    gene;
                                                                            cytostatic;
                                                                                           marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer; amplify
r; PCR; PSA; ss.
                                                                                                                                                                                                                                                                                                                                                                                                          줐
                                                                                                                                                                                                                                           31 G; 27 T; 0 other;
                                                                                                                                                                                                              0
                                                                     SS
                                                                                                                                                                                39
                                                                                                                                                                                                                     Score 24; Pred. No. (
                                                                                                                                                                                                              Mismatches
                                                                                            CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amplify; prostate specific antigen;
                                                                          carcinogen; pharmacodyanamic
                                                                                            7902
                                                                                                                                                                                                                                                                                                                                                                   (PSA) and immunological calibrator in commercia
                                                                                                                                                                                                                     0.17;
                                                                                                                                                                                                                            DB 17;
                                                                                                                                                                                                             0;
                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                     commercial
                                                                                                                                                                                                             0;
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ABV07396
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Best Local S
Matches 24
 16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
18-JUL-2000;
13-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-FEB-2000;
16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
18-JUL-2000;
13-DEC-2000;
                                                                                                                                                                                     WO200160860-A2
                                                                                                                                                                                                                                                       Human; prostate pharmacogenomic
                                                                                                                                                                                                                                                                                                        Human prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            patient; (I) is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of t specification or its complement. (I) is useful for:
(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, uses
                                                                                     17-FEB-2000;
                                                                                                                    20-FEB-2001;
                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                         13-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                          ABV07396;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer in a patient;
(d) assessing the efficacy of a therapy for inhibiting prostate in a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1;
                                                                                                                                                     23-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                           ABV07396 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (e) selecting a composition for inhibiting prostate cancer in a patient;(f) assessing the prostate cell carcinogenic potential of a compound;(g) determining whether prostate cancer has metastazed in a patient;(h) assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 ATGTGGGTCCCGGTTGTCTTCCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATGTGGGTCCCGGTTGTCTTCCTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
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2000US-183319P.
2000US-189862P.
2000US-207454P.
2000US-211314P.
2000US-2119107P.
2000US-255281P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              useful as
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2000US-207454P.
2000US-211314P.
2000US-219007P.
2000US-255281P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP;
                                                                                                                      2001WO-US05171
                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            presence of prostate cancer, stage of
                                                                                                                                                                                                                                                                                        expression marker cDNA 7387.
                                                                                                                                                                                                                                         cancer; cytostatic; carcinogen; marker; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20
                                                                                                                                                                                                                                                                                                                                                                                                           cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ₽.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            a pharmacodyanamic or pharmacogenomic marker
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                                                                                                                                                                                                                                                                                                                                                                                                          ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             G; 43 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Œ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ມສ 23;
. 0.18;
hes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 other;
                                                                                                                                                                                                                                                                       pharmacodyanamic marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 171;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              prostate cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             tate of
useful
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RESULT 11
ABV373300
ID ABV373730
ID ABV37
AC ABV
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                               17-FEB-2000; 2000US-183319P.
16-MAR-2000; 2000US-189862P.
25-MAY-2000; 2000US-207454P.
09-JUN-2000; 2000US-211314P.
18-JUL-2000; 2000US-219007P.
13-DEC-2000; 2000US-255281P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:
(a) assessing whether a patient is affilicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, used for detecting presence of prostate cancer, stage of prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 308 BP; 65 A; 100 C; 79 G; 63 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (e) selecting a composition for inhibiting prostate cancer in a patient; (f) assessing the prostate cell carcinogenic potential of a compound; (g) determining whether prostate cancer has metastasized in a patient; (h) assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 1194; 11750pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; prostate cancer; cytostatic; pharmacogenomic marker; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human prostate expression marker cDNA 37321.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABV37330 standard; cDNA; 428
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                                                         Schlegel R,
                                                                                                                                                                                                                                                                                                                                                                                                                          23-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200160860-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                 20-FEB-2001; 2001WO-US05171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (I) is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       198
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24; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            also useful as a pharmacodyanamic or pharmacogenomic marker
                                                                                                               MILLENNIUM PREDICTIVE MEDICINE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Endege WO,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                         Endege
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                                                            Monahan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 24;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      carcinogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB'23; Length 308;
0.19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pharmacodyanamic marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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The invention relates to an isolated nucleic acid molecule (1) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV062213) of the specification or its complement. (1) is useful for:
(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, used for detecting presence of prostate cancer, stage of prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                             (c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient; (d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient; in a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 7668; 11750pp; English.
                                                                                                                                         Sequence 428 BP; 82 A; 132 C; 123 G; 91 T; 0 other;
                                                                                                                                                                                                       patient;(I) is also useful as a pharmacodyanamic or pharmacogenomic marker.
                                                                                                                                                                                                                                                               (e) selecting a composition for inhibiting prostate cancer in a patient; (f) assessing the prostate cell carcinogenic potential of a compound; (g) determining whether prostate cancer has metastasized in a patient; (h) assessing the aggressiveness or indolence of prostate cancer in a
                                            Similarity
           100.0%; ilarity 100.0%; Conservative 0;
              0;
                                            Pred.
                                                                          Score 24;
Mismatches 0;
                                                   No.
                                            0.19;
                                                                                 DB 23;
                                                                                 Length 428;
                      Indels
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B Best Loc Matches Query Match Local 294

QΥ

RESULT 12 AAA61482 AAA61482 standard; DNA; 517

Prostate specific antigen gene fragment. 23-OCT-2000 PSA gene al DNA amplification; template; prostate sp (first entry) specific nucleic c acid syr antigen; synthesis;

human;

detection;

18-MAY-2000 WO200028082-A1.

09-NOV-1998; 98JP-0317476

08-NOV-1999;

99WO-JP06213

(EIKE ) EIKEN KAGAKU KK

WPI; 2000-376587/32 Notomi T, Hase T;

Nucleic acid synthesis using primer containing a sequence identical to one on the template for isothermal amplification using simple reagents

Example 8 Fig 17; 95pp; Japanese.

CCCCXXXX The invention relates to a novel method of isothermal DNA amplification. A region is produced at the 3' end of a target sequence which contains sub-sequences designated F1, F2C and F1C, where F1C is the complement of F1. A loop is formed by annealing of F1 to its complementary sequence

999999999999988

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RESULT 13
AAH41845
The resent invention describes a new method for synthesising nucleic acids comprising: (a) preparing a nucleic acid, with an F2c region, able to form a loop by annealing the F1 region of the 3' end to the F1c region on the same chain; (b) synthesis of a complementary chain using the 3' end of F1 as the starting point; (c) replacing the complementary chain by annealing an oligonucleotide containing F2, (complementary to F2c) at its 3' end and using it as a starting point for complementary chain synthesis by a polymerase; and (d) replacing the chain synthesised in step c by annealing a polynucleotide containing a region complementary to a part of the chain and using it 3' end as the starting point. Also described are: (1) a kit for synthesising nucleic acids; and (2) a kit for detecting target base sequences. The present sequence represents a prostate specific antigen related nucleotide containing a region is sequence, which is used in an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthesizing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Notomi T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (EIKE ) EIKEN KAGAKU KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Prostate specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Prostate specific antigen related nucleotide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAH41845;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAH41845 standard; DNA; 517
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 517
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3' end
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                                                                                                                                                                                                                                                                                                                                                                                                                        8; Fig 17; 96pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleic acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antigen; nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 24;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           151 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       synthesis; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.2;
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Sequence 517

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94 A; 163 C;

151 G;

109 T; 0 other

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1 ATGTGGGTCCCGGTTGTCTTCCTC 24

Matches

24;

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RESULT 14
ABV25078
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Best Local S
Matches 24
              Query Match
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                                                                                                                                                                           The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;

(b) monitoring the progression of prostate cancer in a patient;

(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                             cancer in a patient;
cancer in a patient;
can seem of a therapy for
                                                                                                                                                                                                                                                                          Novel isolated nucleic acid molecule associated with cancerous prostate cells and correlating with presence of prostate cancer for detecting presence of prostate cancer, stage of prostate cancer.
                                                                                                                                                                                                                                                               Claim 1; Page 4874; 11750pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human
                                                 Sequence
                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                                                                                             Schlegel
                                                                                                                                                                                                                                                                                                                                                                                                                                          16-MAR-2000;
25-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200160860-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; prostate pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABV25078;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABV25078 standard; cDNA; 841
                                                                                   patient;
                                                                                                                     (e) selecting
(f) assessing
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                                                                                                                                                                                                                                                                                                                                                                                                            13-DEC-2000;
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                                                                                             assessing
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                                                                       also useful as a pharmacodyanamic or pharmacogenomic
              Similarity
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                                                 841 BP;
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2000US-219007P.
2000US-255281P.
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                                                                                                                                                                                                                                                                                                                                                             Endege WO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001WO-US05171
                                                                                           a composition for inhibiting prostate cancer in a patient; the prostate cell carcinogenic potential of a compound; ng whether prostate cancer has metastasized in a patient; the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                             2000US-189862P
2000US-207454P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     marker;
                                                  144
                                                                                                                                                                                                                                                                                                                                                                                  PREDICTIVE MEDICINE INC.
              100.0%;
                                               A; 262 C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cytostatic;
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Score 24; DB 23, Pred. No. 0.21;
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Pred. No. .0.
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                                              262 G; 173 T; 0 other;
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                                                                                                                                                      inhibiting prostate cancer
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                                                                                                                                                                                                                                                                                      prostate cancer, useful
e of prostate cancer -
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                                                                        marker
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Search completed: November 19, 2002, 03:06:36 Job time: 171.462 secs
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                                                                                  Ouery Match 100.0%; Score 24; DB 21; Length 990; Best Local Similarity 100.0%; Pred. No. 0:21; Matches 24; Conservative 0; Mismatches 0; Indels
                                                                                                                                               The invention relates to a method of immunization against prostatic cancer (PC) by administering a nucleic acid (this sequence) that encodes prostate-specific antigen (PSA), or its immunologically active fragment, cells and then expressed to generate an immune response against PSA. The method is used to protect against, or treat, PC, or more generally to induce a cellular immune response to PSA;
                                                                                                                              Sequence 990 BP; 209 A; 291 C; 273 G; 217 T; 0 other;
                                                                                                                                                                                                                                                    Disclosure; Page 42; 47pp; English.
                                                                                                                                                                                                                                                                    Immunization against prostatic cancer using nucleic acid encoding prostate-specific antigen, for treatment or prophylaxis -
                                                                                                                                                                                                                                                                                                              WPI; 2000-072551/06.
P-PSDB; AAY56048.
                                                                                                                                                                                                                                                                                                                                               Weiner DB, Kim JJ;
                                                                                                                                                                                                                                                                                                                                                                     (UYPE-) UNIV PENNSYLVANIA.
                                           29-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                     28-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    W09961068-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunization; prostate cancer; prostate-specific antigen; PSA;
cytostatic; immune response; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human prostate-specific antigen gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-MAR-2000 (first entry)
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Post-processing: Minimum Match 0%
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                                                                                                                                                                                                                                                                                                                                                            seq length: 0 seq length: 2000000000
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24
November 19, 2002, 01:43:19;
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Copyright (c) 1993 - 2002 Compugen Ltd
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283.494 Million cell updates/sec
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Pred. No. and is No. is the number of results predicted by chance to have a greater than or equal to the score of the result being printed, s derived by analysis of the total score distribution.

## SUMMARIES

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Result		Query				*
No.	Score	Match	Match Length DB	BB	ID	Description
1	24	100.0	316	و	100.0 316 9 AA593245	AA593245 nn07e10.s
N	24	100.0	336	9	AA506459	AA506459 nh46a08.s
w	24	100.0	371	9	AA503943	AA503943 nh38d05.s
c 4	24	100.0	391	9	AA506939	AA506939 nh64b07.s
υ	24	100.0	408	12	BF678836	BF678836 602153143
σ	24	100.0		9 A	AA528287	AA528287 nh26d08.s

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22 24 24	222222	266666 266666 266666666666666666666666	2222222	2222222	222222222222222222222222222222222222222
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RESULT 1
AA593245
LOCUS
DEFINITION

SOURCE ORGANISM ACCESSION VERSION KEYWORDS

AA593245.1 GI:2409007 EST.

Homo sapiens

human

mRNA sequence.

AA593245

nn07e10.s1 NCI\_CGAP\_Pr4.1 Homo sapiens similar to gb:M21895 PROSTATE SPECIFIC

316 bp

mRNA

A linear EST 25-SEP-1997 cDNA clone IMAGE:1077066 ANTIGEN PRECURSOR (HUMAN);

ALIGNMENTS

TITLE AUTHORS NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicyap. National Cancer Institute, Cancer Genome Anatomy Project (CGAP), Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 316)

REFERENCE

JOURNAL

COMMENT

Tumor Gene Index Unpublished (1997)

Contact: Robert Strausberg, Ph.D.

Email: cgapbs r@mall.nih.gov

Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.,

Michael R. Emmert-Buck, M.D., Ph.D.

CDNA Library Preparation: David B: Krizman, Ph.D.

cDNA Library Preparation: David B: Krizman, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

Insert Length: 427 Std Error: 0.00

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AUTHORS
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

El (bases 1 to 336)

RS NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
Tumor Gene Index
Unpublished (1997)

Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov

Tissue Procurement: David G. Bostwick, M.D., Rodrigo F. Chuaqui,
M.D., Michael R. Emmert-Buck, M.D., Ph.D.
CDNA Library Preparation: David B. Krizman, Ph.D.
CDNA Library Preparation: David B. Krizman, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LINL at:
www-bio.llnl.gov/Dbrp/image/image.html
Insert Length: 442

Std Error: 0.00

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TOTATION/Onalifiare
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High quality sequence stop: 293.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="NCI_CGAP_Pr4.1"
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/db_xref="taxon:9606"
/clone="IMAGE:1077066"
                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:955382"
/clone_lib="NCI_CGAP_Pr5"
                 /sex="male"
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/lab_host="DH10B"
                                                                                                                                                                   Location/Qualifiers
/note="Vector: pAMP10; mRNA made from normal prostatic
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Contact: Robert Strausberg, Ph.D.
Email: Gapbs-remail.nih.gov
Tissue Procurement: David G. Bostwick, M.D., Rodrigo F. Chuaqui,
M.D., Michael R. Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: David B. Krizman, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-GAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
www-bb.llnl.gov/bbrp/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inage/inag
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nh384005.s1 NCI_CGAP_PT5 Homo sapiens cDNA clone IMAGE:954633
similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);,
mRNA sequence.
AA503943 77770077
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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/clone="tib="NCI_CGAP_Pr5"
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/tissue_type="prostate"
/lab_host="DHIOB"
/lab_host="DHIOB"
/note="vector: pawe10; mRNA made from normal prostatic printelial cells, cDNA made by oligo-dT priming.
won-directionally cloned. Size-selected on agarose gel, average insert size 600 bp.
a verage insert size 600 bp. "
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Email: cgapbs r@mail.nih.gov

Tissue Procurement: David G. Bostwick, M.D., Rodrigo F. Chuaqui,
M.D., Michael R. Emmert-Buck, M.D., Ph.D.

cDNA Library Preparation: David B. Krizman, Ph.D.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 488 Std Error: 0.00

Seq primer: -40m13 fwd. ET from Amersham.
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391 bp mRNA linear EST 20-AUG-1997 nh64b07.sl NCI_CGAP_Pr8 Homo sapiens cDNA clone IMAGE:957109 similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);
                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mRNA sequence.
BF678836
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    602153143F1 NIH_MGC_83 Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note-"Vector: pamP10; mRNA made from invasive prostate tumor, cDNA made by oligo-dT priming. Non-directionally cloned. Size-selected on agarose gel, average insert size 600 bp. "
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/db_xref="taxon:9606"
/clone="IMAGE:957109"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /tissue_type="prostate"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="NCI_CGAP_Pr8"
/sex="male"
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1.9;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE
AUTHORS
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ORGANISM
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AA528287
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ORIGIN
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                                                                                                                                                                                     source
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37 ATGTGGGTCCCGGTTGTCTTCCTC 60
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                                                                                                                                                                                                                                                                                                                                                              Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.,
Michael Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA528287

A57 bp __mrNA linear EST 20-AUG-1997

nh26d08.sl NCI_CGAP_Pr3 Homo sapiens cDNA clone IMAGE:953487

similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 457)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                  found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html seq primer: -40ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                 High quality sequence stop: 187.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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                                                                                                                                                                                                                                                   CDNA Library Preparation: David, B. Krizman, Ph.D.
CDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /incte-Torgan: prostate; Vector: pDNN-LIB (Clontech);
/note-Torgan: prostate; Vector: pDNN-LIB (ggccattatggcc);
Site_1: SfiI (ggccgcctcggcc); Site_2: SfiI (ggccattatggcc);
5' and 3' adaptors were used in cloning as follows: 5'
adaptor sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor
sequence: 5'-ATTCTAPAGGCCGAGCGGCGCACATG-dT(30)BN-3'
(where B = A, C, or G and N = A, C, G, or T). Average
insert size 1.4 kb (range 0.5-4.0 kb). 14/15 colonies
contained inserts by PCR. This library was enriched for
full-length clones and was constructed by Clontech
Laboratories (Palo Alto, CA)."

74 a 127 c 115 g 92 t
/organism="Homo sapiens"
/db_xref="taxon:9606"
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/db_xref="taxon:9606"
/clone="IMAGE:4294258"
/clone_lib="NIH_MGC_83"
                                                                                             location/Qualifiers
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/clone\_lib="NCI\_CGAP\_Pr3"

/clone="IMAGE:953487"

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1 ATGTGGGTCCCGGTTGTCTTCCTC 24
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Butheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 465)
Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,
Dias Neto,E., Garcia Correa,R., Bordin,S., Costa,F.F.,
Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,
Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare
,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 465 bp mRNA linear EST 14-JUN-200 MR3-FN0209-070201-010-b11 FN0209 Homo sapiens cDNA, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010,
                                                                                                                                                                                                                                                                                                                                                                                          Dimail: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=MR36t2=MR3-FN0209-
070201-010-bilstj=2001-02-078t4=1)
Seq_primer: puc 18 forward
seq_primer: puc 18 forward
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                   quality sequence start: 7 quality sequence stop: 465.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Natl. Acad. Sci. U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /dev_stage="45 years old"
/lab_host="DH10B"
   /note="Organ: prostate_normal; Vector: puc18; Site_1: SmaI; site_2: SmaI; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse
Research) profiles into
                                                                                                                                                    /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone_lib="FN0209"
/dev_stage="Adult"
                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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Pred. No. 1.9;
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REFERENCE AUTHORS

COMMENT

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JOURNAL MEDLINE

RESULT 7 BI046475 LOCUS

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26

ACCESSION VERSION

DEFINITION KEYWORDS

SOURCE ORGANISM

BASE COUNT ORIGIN

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KEYWORDS
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutherla; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 495)
Dias Neto, E. Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R.,
Magai, M.A., da Silva-M., Jr., Zago, M.A., Bordin, S., Costa, F.F.,
Goldman, G.H., Carvalho, A.F., Matsukuma, A., Baid, G.S., Simpson, D.H.,
Brunstein, A., deoliveira, P.S., Bucher; P., Jongeneel, C.V., O'Hare,
M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BE840533 495 bp mRI RC1-FN0188-260700-022-f06 FN0188 Homo BE840533
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24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                      bmail: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scritpts/gethtml2.pl?tl=&t2=RC1-FN0188-260
700-022-f06&t3-200-07-26&t4-1)
Seq primer: puc 18 forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tel: +55-11-2704922
Fax: +55-11-2707001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Simpson A.J.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence tags
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                                                                                             Similarity
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                                                                              Conservative
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a 134 c 144 g 103 t 1 others
                                                                                                                                                                /note="Organ: prostate_normal; vector: puc18; Site_1: Smal | Site_2: Smal; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Pattent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
                                                                                                                                                                                                                                                                                                                      /db_xref="taxon:9606"
/clone_lib="FN0188"
/dev_stage="Adult"
                                                                                                                                                                                                                                                                                                                                                                            /organism="Homo sapiens"
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Dias Neto, E., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R., Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F., Goldman, G.H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H., Brunstein, A., deOliveira, P.S., Bucher, P., Jongeneel, C.V., O'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and
                                                                              np08f03.s1 NCI_CGAP_Pr3 Homo s
similar to gb:M21895 PROSTATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
                                       mRNA sequence.
AA639901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
AA639901.1 GI:2563680
EST.
                                                                                                                       AA639901
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (http://www.ludwig.org.br/scripts/gethtml2.pl?tl=&t2=RC1-FN0188-210
700-021-a02&t3=2000-07-21&t4=1)
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                                                                                                                                                                                                                                                                                                                                                                                             /note-"Organ: prostate_normal; Vector: puc18; Site_1: SmaI; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /db_xref="taxon:9606"
/clone_lib="FN0188"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /dev_stage="Adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
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Pred. No.
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                                                                                                                     523 bp
                                                                      איא EST 23-OCT-1997 sapiens cDNA clone IMAGE:1115741
SPECIFIC ANTIGEN PRECURSOR (HUMAN);,
                                                                                                                                                                                                                                                                                                                     DB 12;
1.9;
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Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.,
Michael Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: David, B. Krizman, Ph.D.
cDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-GGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                      Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvelho,A.F., Matsukuma,A., Baia,G.S., Simpson,Brunstein,A., deOllveira,P.S., Bucher,P., Jongeneel,C.V., O'Ha,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                           Shotgun sequencing of the human transcriptome with ORF expressed
                                                                     Simpson, A.J.
                                                                                                                                                                                    Dias Neto, E.,
                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                EST
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1 (bases 1 to 523)
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Mammalia; Eutheria; Primates; Cațărrhini; Hominidae;
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Seq primer: -40ml3 fwd. ET from Amersham
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                                                                                                                                                                          (bases 1 to 552)
as Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="vector: pAMP10; Site_1: Not1; Site_2: ECORI; 1st strand cDNA was primed with oligo(dT)17 on 50 ng of DNAse-treated, total cellular RNA obtained from 5,000-10,000 microdissected cells histologically-determined to be fully malignant prostate cancer cells. Double-stranded cDNA was ligated to EcoRI adaptors, 5 cycles of PCR applied to the cDNA with an adaptor-specific primer, and the resulting PCR product subcloned into pAMP10 by the UDG-cloning method (Life Technologies). Average insert size is 600 bp. NOTE: Not directionally cloned. This library was constructed by David Krizman."
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/lab_host="DH10B"
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/db_xref="taxon:9606"
/clone="IMAGE:1115741"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="NCI_CGAP_Pr3"
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Acad. Sci. U.S.A.
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Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,
Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Soldman,G.H., Carvalho,A.F., Matsukuma,A., Bala,G.S., Simpson,D.H.,
Brunstein,A., deoliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare
M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=st2=RCl-FN0188-140
800-011-g05st3=200-08-14st4-1)
seq primer: puc 18 forward
High quality sequence start: 2
High quality sequence start: 2
Location/Qualifiers
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BE840537
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Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 and
Fax: +55-11-2707001
Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo:
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                                                                                                                         Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 and
                                                                                                                                                                                      Contact: Simpson A.J.G.
                                                                                                                                                                                                                                                                       Shotgun sequencing of the human transcriptome with ORF expressed
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                                                                                +55-11-2704922
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//db_xref="taxon:9666"
//db_xref="taxon:9666"
//clone_ilb="FxN0188"
//dev_stage="Adult"
//note="Organ: prostate_normal; Vector: puc18; Site_1: SmaI; Site_2: SmaI; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
79 a 150 c 171 g 152 t
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540 ATGTGGGTCCCGGTTGTCTTCCTC 517
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: cgapbs-remail.nih.gov
Tissue Procurement: CLONETECH Laboratories, Inc.
cDNA Library Preparation: CLONETECH Laboratories, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM1145 row: a column: 10
High quality sequence stop: 587.
Location/Qualifiers
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700-022-h06&t3=2000-07-26&t4=1)
Seq primer: puc 18 forward
High quality sequence start: 10
High quality sequence stop: 572.
Location/Qualifiers
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NIH-MGC http://mgc.nci.nih.gov/...
National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished (1999)

Contact: Robert Strausberg, Ph.D.
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/db_xref="taxon:9666"
/clone_jlb="FxN018B"
/clone_jlb="FxN018B"
/dev_stage="Adult"
/dev_stage="Adult"
/note="Organ: prostate_normal; Vector: puc18; Site_1: SmaI
/Site_2: SmaI, A mini-library was made by cloning
products derived from ORESTES PCR (U.S. Letters Patent
application No. 196,716 ' Ludwig Institute for Cancer
Research) profiles into the pUC 18 vector. Reverse
transcription of tissue.mRNA and cDNA amplification were
performed under low stringency conditions."
56 a 175 c 156 g 85 t
/ Organisma*Homo sapiens // Organisma*Homo sapiens // Organisma*Homo sapiens // Organisma*Homo sapiens // Clone="IMAGE:4294929" // Clone="IMAGE:4294929" // Clone="Iba*NIH_MGC_83" // Note="Organ: prostate; Vector: pDNR-LIB (Clontech); // Note="Organ: prostate; Vector: pDNR-LIB (Clontech); // Site_1: Sfii (ggccgctctggcc); Site_2: Sfii (ggccattatggcc); Site_1: Sfii (ggccgctctggcc); South cloning as follows: 5', and 3' adaptors were used in cloning as follows: 5', adaptor sequence: 5'-ATTCTAGAGGCCGATTATGGCC-3' and 3' adaptor sequence: 5'-ATTCTAGAGGCCGAGGGCGGCGCAATG-GT(30)BN-3' (where B = A, C, or G and N = A, C, G, or T). Average insert size 1.4 kb (range 0.5-4.0 kb). 14/15 colonies contained inserts by PGR. This library was enriched for full-length clones and was constructed by Clontech
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RESULT 15
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                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
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                                                                                                                                                                                                                        1 ATGTGGGTCCCGGTTGTCTTCCTC 24
                                                                                                                                                                                        ATGTGGGTCCCGGTTGTCTTCCTC 66
BF677605 648 bp mRNA linear EST 21-DEC-2000 602085679F1 NIH_MGC_83 Homo sapiens cDNA clone IMAGE:4249939 5',
                                                                                                                                                                                                                                                                                                                   24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: cgapbs-r@mail.nih.gov
Tissue Procurement: CLONETECH Laboratories, Inc.
cDNA Library Preparation: CLONETECH Laboratories, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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/db_xref="taxon:9606"
/clone="IMAGE:4295164"
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/note="Organ: polyner; Site_2: Sill (ggccattatggcc);
/note="Organ: polyner; Sill (ggccattatggcc);
/not
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Search completed: November 19, Job time: 1377.08 secs

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REFERENCE
AUTHORS
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JOURNAL
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SOURCE
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VERSION
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43 ATGTGGGTCCCGGTTGTCTTCCTC
                     1 ATGTGGGTCCCGGTTGTCTTCCTC 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Robert Strausberg, Ph.D.: 
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: CLONETECH Laboratories, Inc.
cDNA Library Preparation: CLONETECH Laboratories, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov plate: L/LML070 row: n column: 20 High quality sequence stop: 611.
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National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Clone distribution: MGC clone distribution information can be
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                                                                                        Conservative
                                                                                                                                                                                                  /note-"organ: prostate; Vector: pDNR*LIB (Clontech);
Site_1: Sfil (ggccgctcggcc); Site_2: Sfil (ggccattatggcc);
5' and 3' adaptors were used in cloning as follows: 5'
adaptor sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor
sequence: 5'-ATTCTAGAGGCCGAGGCGGCCATG-GT(30)BN-3'
(Where B = A, C, Or G and N = A, C, G, Or T). Average
insert size 1.4 kb (range 0.5-4.0 kb). 14/15 colonies
contained inserts by PGR. This library was enriched for
full-length clones and was constructed by Clontech
Laboratories (Palo Alto, CA)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                             /clone="IMAGE:4249939"
/clone_lib="NIH_MGC_83"
/lab_host="DH10B (T1 phage-resistant)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /db_xref="taxon:9606"
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Maximum Match 100%
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Perfect score:
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1: /cgn2_6/ptodata/2/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/2/ina/5B_COMB.seq:*
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5: /cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*
5: /cgn2_6/ptodata/2/ina/Dackfiles1.seq:*
        IDENTITY_NUC
Gapop 10.0 , Gapext 1.0
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US-09-210-016-2
US-08-358-782D-13
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US-08-358-782D-14
US-08-358-782D-14
US-08-605-785-607-
US-08-605-785-607-
US-08-605-785-607-
US-08-90-09-630-31
US-09-09-042-071-19
US-08-454-720A-38
US-08-983-975D-10
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US-08-454-720A-42  Sequence 42, Application US/08454720A  Patent NO. 576688  GENERAL INFORMATION:  APPLICANT: Sobol, Robert E.  APPLICANT: Green, Mark R.  APPLICANT: Kawasaki, Ernest S.  TITLE OF INVENTION: Detection of Carcinoma Metastases by TITLE OF INVENTION: Nucleic Acid Amplification  NUMBER OF SEQUENCES: 42  CORRESPONDENCE ADDRESS:  ADDRESSEE: Hoffman-La Roche Inc.  STREET: 340 Kingsland Street  CITY: Nutley  STATE: New Jersey  COUNTRY: U.S.A.	Query Match Best Local Similarity 100.0%; Score 24; DB 4; Length 70; Best Local Similarity 100.0%; Pred. No. 0.022; Matches 24; Conservative 0; Mismatches 0; Indels 0; Gaps  Qy 1 ATGTGGGTTCCGGTTGTCTTCCTC 24	nalized Microspheres fosensors	28 16 66.7 871 4 US-08-969-987-7 29 16 66.7 1279 3 US-08-985-950-5 30 16 66.7 1341 4 US-08-985-950-5 31 16 66.7 1344 4 US-08-983-075D-6 32 15.8 66.7 1728 3 US-08-983-075D-8 33 15.8 65.8 424 1 US-08-158-189-1 34 15.8 65.8 424 1 US-08-158-189-1 35 15.6 65.0 340 4 US-08-158-189-1 36 37 15.6 65.0 1508 4 US-08-158-199-1 37 15.6 65.0 1508 4 US-09-404-390-14 38 15.6 65.0 2333 4 US-09-404-390-14 39 15.2 63.3 1512 2 US-08-576-6,22A-2 39 15.2 63.3 1512 2 US-08-898-219B-20 39 15.2 63.3 1512 2 US-08-898-219B-20 39 15.2 63.3 2241 3 US-09-233-336A-20 39 15.2 63.3 2241 3 US-09-233-336A-20 39 15.2 63.3 2241 4 US-09-2036-20 39 15.2 63.3 2241 4 US-09-204-226-20 39 15.2 63.3 2241 4 US-09-204-226-20 39 15.2 63.3 2241 4 US-09-904-226-20
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US-08-358-782D-13
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INFORMATION FOR SEQ ID NO: 42
SEQUENCE CHARACTERISTICS:
LENGTH: 111 base pairs
TYPE: nucleic acid
STRANDENNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.01, version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/454.720A
FILING DATE: May 31, 1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNDER: 08/357,565
PILING DATE: December 16, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Petry, Douglas
REGISTRATION NUMBER: 35321
REGISTRATION NUMBER: 35321
REGISTRATION NUMBER: 35321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 100.0%; Score 24; Best Local Similarity 100.0%; Pred. No. Matches 24; Conservative 0; Mismatch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1
Patent No.
APPLICATION NUMBER: US/08/358
FILING DATE: 15-DEC-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Beardell, Lori Y.
REGISTRATION NUMBER: 34.293
REFERENCE/DOCKET NUMBER: TJU-
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Gorec, Carlo
APPLICANT: Gomella, Leonard
APPLICANT: Mulholland, S. Grant
APPLICANT: Mulholland, S. Grant
APPLICANT: Mulholland, S. Grant
APPLICANT: Musholland, S. Grant
APPLICANT: Fischer, Rainer
TITLE OF INVENTION: Methods of Detecting Micrometastasis of
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz &
STREET: One Liberty Place 46th. Floor
                                                                                                                                                                                                                               STATE: PA
ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBÉR: 35321
REFERENCE/DOCKET NUMBER: 9178
REFERENCE/DOCKET NUMBER: 9178
RELEPHONE: (510) 814-2974
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                                                            TJU-1327
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.023;
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1 ATGTGGGTCCCGGTTGTCTTCCTC 24

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HYPOLOGY: linear MOLECULE TYPE: DNA (
HYPOTHETICAL: NO ANTI-SENSE: NO US-08-764-527A-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TOPOLOGY: linear
; MOLECULE TYPE: DNA
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
US-08-358-782D-13
                                                                                                                                                        PILING DATE:
CLASSIFICATION:
CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: US/08/358,782
APPLICATION NUMBER: US/08/358,782
ATTORNEY/AGENT INFORMATION:
NAME: Beardell, Lori Y.
REGISTRATION NUMBER: 34,293
REFERENCE/DOCKET NUMBER: TJU-1327
TELEPHONE: 215-568-3100
TELEPHONE: 215-568-3100
TELEPAX: 215-568-3439
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 992 base pairs
TYPE: nucleic acid
TTYPE: nucleic acid
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US-08-764-527A-13
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  Query Match
Best Local S
Matches 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/764,527A
FILING DATE:
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LENGTH: 992 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Croce, Carlo
APPLICANT: Gomella, Leonard
APPLICANT: Mulholland, S. Grant
APPLICANT: Moreno, Jose
APPLICANT: Fischer, Rainer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Methods of Detecting Micrometastasis of TITLE OF INVENTION: Prostate
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STATE: PA
STATE: 19103
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h 100.0%; Score 24; DB 2; Similarity 100.0%; Pred. No. 0.028; 24; Conservative 0; Mismatches
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Fischer, Rainer
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                                                                                                                                                                                   tent No.
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LENGTH: 1462 base pairs
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CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
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                                                                                                                                                                INFORMATION:
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24; Conserv
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Beardell, Lori Y.
Beardell, 101 Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCES:
                                                                                                                                                                                                  Application US/08764527A
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                                                                               Fischer, Rainer
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llarity 100.0%;
Conservative (
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                                                                                                Moreno, Jose
                                                                                                              Mulholland, S. Grant
                                                                                                                                 Gomella, Leonard
                                                                                                                                                  Croce, Carlo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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Woodcock Washburn Kurtz Mackiewicz &
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                                                               Methods of Detecting Micrometastasis
                                                Prostate
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Pred. No. 0
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                                                                                                        SEQ ID NO 31
LENGTH: 7130
TYPE: DNA
ORGANISM: Homo sapiens
                               Matches
                                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                Sequence 31, Appli Patent No. 6287569
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                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/056,105
CURRENT FILING DATE: 1998-04-06
                                                                                                                                                                                       EARLIER FILING DATE: 1997-04-10 NUMBER OF SEQ ID NOS: 35
                                                                                                                                                                                                                       CURRENT FILING DATE: 1998-04-06 EARLIER APPLICATION NUMBER: 60/043,467
                                                                                                                                                                                                                                                                     FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                APPLICANT: KIPPS, THO APPLICANT: WU, YUNQI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 215-568-3439
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                TITLE OF INVENTION: VACCINES WITH ENHANCED INTRACELLULAR TITLE OF INVENTION: PROCESSING
                                                                                                                                                                            SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: TJ
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 15-DEC-1994 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              25 ATGTGGGTCCCGGTTGTCTTCCTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE:
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1 ATGTGGGTCCCGGTTGTCTTCCTC 24
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                               l Similarity
24; Conserv
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                                                                                                                                                                          FastSEQ for Windows Version 3.0
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                               Conservative
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                                            100.0%; Score 24; I
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                             Mismatches
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                                              .032;
                                                           4; Length 7130;
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                             Gaps
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US-09-605-785-607

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APPLICANT: Skeiky Yasir A.W.

APPLICANT: Hepler, William

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND FILE REFERENCE: 210121.427C16

CURRENT APPLICATION NUMBER: US/09/605,785

CURRENT ETLING DATE: 2000-06-27

NUMBER OF SEQ ID NOS: 835

SOFTWARE: FASTED for Windows Version 3.0

SEO ID NO 607

LENGTH: 22

TYPE: NOT
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Patent No. 5614372
GENERAL INFORMATION:
APPLICANT: Lilja, Hans
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Best Local 9
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .PPLICANT: Xu, Jiangchun
.PPLICANT: Dillon, Davi
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/394,033
                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Venable, Baetjer, Howard & Civiletti
STREET: 1201 New York Avenue, NW, Suite 1000
                                                                                                                                                                                                                                                     APPLICANT: Lilja, Hans
APPLICANT: Lundwall, Ake
APPLICANT: Lundwall, Ake
APPLICANT: Lowgren, Janita
TITLE OF INVENTION: Early Detection of Prostate Cancer (CAP)
TITLE OF INVENTION: by Employing Prostate Specific Antigen (PSA) and Human
TITLE OF INVENTION: clandular Kallikrein (HGK-1)
                                                                                                                                                             ADDRESSEE: Venabli
STREET: 1201 New
CITY: Washington
STATE: DC
                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 ATGTGGGTCCCGGTTGTCTT 20
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20; Conserv
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Carter, Darrick
Li, Samuel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fanger, Gary R. Retter, Marc W. Stolk, John A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Henderson, Rob
Kalos, Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dillon, Davin C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Harlocker,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mitcham, Jennifer L.
                                                                                                                                             USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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ick, Thomas S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  83.3%; Scc
100.0%; Pr
''' 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 20; DB 4; Length 22; pred. No. 1.3; 0; Mismatches 0; Indels
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; MOLECULE TYPE: cl;
HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE:
GRANISM: HOMO
US-08-394-033-1
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APPLICATION NUMBER: GB 9512996.1 ETLING DATE: 26-JUN-1995
ATTORNEY/AGENT INFORMATION:
NAME: Cherry, David A.
REGISTRATION NUMBER: 35,099
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
TELEPAX: (215) 568-3139
INFORMATION FOR SEQ ID NO: 31:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 10
US-08-577-492-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                      COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC DOS/MS-DOS
SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/577,492
FILING DATE: 22-DEC-1995
CLASSIFICATION UNMBER: GB 9426227.6
APPLICATION NUMBER: GB 9426227.6
FILING DATE: 23-DEC-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 951296.1
APPLICATION NUMBER: GB 951296.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 202-962-8300
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Perry, Martin John
APPLICANT: Lumb, Simon Mark
TITLE OF INVENTION: HUMAN PHOSPHODIESTERASE TYPE IVC, AND
TITLE OF INVENTION: ITS PRODUCTION AND:USE
NUMBER OF SEQUENCES: 40
CORRESPONDENCE ADDRESS:
ADDRESSE: Woodcock, Washburn, Kurtz, Mackiewicz & No.
STREET: One Liberty Place, 46th floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Innen, Jeffrey L.
NAME: Innen, Jeffrey L.
REGISTRATION NUMBER: 28,957
REFERENCE/DOCKET NUMBER: 23
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-62-4810
TELEPHONE: 202-62-8200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 28 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: UZIP: 19103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: One LINGLY CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 2:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 75.0%; Score 18; DB 1; Length 28; Local Similarity 100.0%; Pred No. 10; Local Similarity 0.00 No. 10; October 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1, Application US/08577492 5851784
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         USA
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N: 435
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Martin John
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SEQUENCE CHARACTERISTICS:

TYPE: nucleic acid STRANDEDNESS: double

TOPOLOGY:

linear

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US-09-079-630-31
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                                                                                                                             US-09-079-630-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 31, Application Patent No. 6291199
GENERAL INFORMATION:
                                                              Matches
                                                                                            Query Match
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                                                                                 Best
                                                                                                                                                                                       TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 2153 base pairs
                                                                                                                                                                                                                                                                                                                                                FILING DATE: 23-DEC-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9426227.6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                      NAME: Cherry, David A.
REGISTRATION NUMBER: 35,099
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
                                                                                                                                                                                                                                                                                                                     FILING DATE: 26-JUN-1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/577,492
FILING DATE: 22-DEC-1995
APPLICATION NUMBER: GB 9426227.6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: HUMAN PHOSPHODIESTE TITLE OF INVENTION: ITS PRODUCTION AND NUMBER OF SEQUENCES: 40-CORRESPONDENCE ADDRESS:
198 GGTCCCGGTTGTCTTCCT 215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 198 GGTCCCGGTTGTCTTCCT 215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/079,630 FILING DATE:
                                                                               Local
                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: doub
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
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                               6 GGTCCCGGTTGTCTTCCT 23
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al Similarity 100.0%;
18; Conservat;
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                                                                             Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E: Woodcock, Washburn, Kurtz, Mackiewicz & No. 6291199ris
One Liberty Place, 46th floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Perry, Martin John
Lumb, Simon Mark
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                                                                                                                                            linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Floppy disk
                                                                                                                                                           double
                                                                             75.0%;
100.0%;
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                                                              0
                                                                               Score 18; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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Pred. No.
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                                                              Mismatches
                                                                                 15,
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                                                                                            Length 2153;
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                                                              Indels
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RESULT 12

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                                                                                                                                                                                                                                                                                                                                                            Sequence 38, Application US/08454720A Patent No. 5766888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 73.3%;
Best Local Similarity 83.3%;
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/042,071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (206) 682-603
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1548 base pair:
                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                               APPLICANT: Sobol, Robert E.
APPLICANT: Green, Mark R.
APPLICANT: Kawasaki, Ernest S.
TITLE OF INVENTION: Detection of Carcinoma Meta
TITLE OF INVENTION: Nucleic Acid Amplification
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hoffman-La Roche Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09
FILING DATE: 13-MAR-1998
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: KAY, WILLIAM W.
TITLE OF INVENTION: REPLICATION: SMALL OF INVENTION: CONTROL
                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               411 ATGTGGCTTCCGGTTCTCTCCGC 388
                                                                                                                                               STREET: 340 F
       COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                          STATE: New Jersey COUNTRY: U.S.A. ZIP: 07110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 660081.407
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 ATGTGGGTCCCGGTTGTCTTCCTC 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleic acid
DEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6294372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         McMasters, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/09042071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E: SEED and BERRY LLP 6300 Columbia Center,
                                                                                                                                                                   340 Kingsland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Burian, Jan
             IBM PC compatible SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
PatentIn Release #1.01,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            682-6031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REPLICATION GENES AND GENE PRODUCTS FROM SMALL CRYPTIC PLASMIDS AND METHODS FOR CONSTRUCTING CONTROLLED-REPLICATION PLASMID VECTORS
                                                                                                                                                                                                                                                                    Detection of Carcinoma Metastases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ö.
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Pred. No. 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 4; Length 1548;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:

08/357,565 ber 16, 1994

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/454,720A
FILING DATE: May 31, 1995
CLASSIFICATION: 435

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FILING DATE: December 16, 1994
ATTORNEY/AGENT INFORMATION:
NAME: PELTY, DOUGHAS
REGISTRATION NUMBER: 9178
REFERENCE/DOCKET NUMBER: 9178
TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 814-2977
INFORMATION FOR SEQ ID NO: 38:
SEQUENCE CHARACTERISTICS:
LENGTH: 1864 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
MOLECULE TYPE: DNA (genomic)
US-08-454-720A-38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 10, Application US/08983075D

Sequence 10, Application US/08983075D

Patent No. 630361

GENERAL INFORMATION:

APPLICANT: VIHKO, Pirkko

TITLE OF INVENTION: HUMAN GLANDULAR KALLIKREIN-1 (HK2)

FILE REFERENCE: 1491/44025

CURRENT APPLICATION NUMBER: US/08/983,075D

CURRENT FILING DATE: 1998-06-08

PRIOR APPLICATION NUMBER: PCT/F196/00382

PRIOR FILING DATE: 1996-06-28

NUMBER OF SEQ ID NOS: 14

SOFTWARE: Patentin version 3.0

SEQ ID NO 10

LENCTH: 72
                                                                                                                                                                                                                                                                                                                                                        ; NAME/KEY: misc_feature
OTHER INFORMATION: nucleotide sequence encoding the prepro-sequence
OTHER INFORMATION: of the hK2 protein
US-08-983-075D-10
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US-08-983-075D-10
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                                                                                                                     RESULT 15
US-09-385-982-435/c
Sequence 435, Application US/09385982
Patent No. 6262334
GENERAL INFORMATION:
APPLICANT: ENDEGE, WILSON O., ET AL.
APPLICANT: ENDEGE, WILSON O., ET AL.
TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION
TITLE OF INVENTION: PRODUCTS: II
                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 68.3
Best Local Similarity 94.4
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Homo sapiens
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1189 GTCCCGGTTGTCCTCCTC 1172
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                                                                                                                                                                                                                                                                                                  66.7%;
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Pred. No. 89;
0; Mismatches
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CURRENT APPLICATION NUMBER: US/09/385,982
CURRENT FILING DATE: 1999-08-30
EARLIER APPLICATION NUMBER: 09/328.111
EARLIER APPLICATION NUMBER: 09/328.111
EARLIER FILING DATE: 1999-06-08
EARLIER FILING DATE: 1999-06-08
EARLIER APPLICATION NUMBER: 60/117,393
EARLIER FILING DATE: 1999-01-27
EARLIER FILING DATE: 1999-01-27
EARLIER FILING DATE: 1998-08-31
NUMBER: OF SEQ ID NOS: 544
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 435
SEQ ID NO 435
LENGTH: 558
TYPE: DNA
ORGANISM: HOMO SAPIENS
FEATURE:
ENAUTY VEV. misc feature
                                                                                                                                                                                                                                                                        ; NAME/KEY: misc_feature
; LCCATION: (1)...(558)
; OTHER INFORMATION: n - A,T,C or G
US-09-385-982-435
                                                                                                                                    QΥ
Search completed: November 19, 2002, 05:20:20 Job time: 54.0769 secs
                                                                                            Вb
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Best Local Similarity 79.3
Matches 19; Conservative
                                                                                            237 ATTIGGGTCCACGCTCTCCTC 214
                                                                                                                       1 ATGTGGGTCCCGGTTGTCTTCCTC 24
                                                                                                                                                                                                      66.7%;
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Gaps

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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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                                                                 nucleic search, using sw model
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! '_cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
! '_cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
! '_cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
! '_cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
! '_cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
! '_cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
! '_cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
! '_cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
! '_cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
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! '_cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
! '_cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
! '_cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
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! '_cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
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110 US-09-755-100-2
110 US-09-755-100-2
110 US-09-759-143-607
110 US-09-822-827-607
110 US-09-974-300-2790
110 US-09-974-300-2790
110 US-09-976-868-429
110 US-09-936-613-4
110 US-09-938-8422-77
110 US-09-938-8428-7207-3
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US-09-755-100-4
US-09-755-100-5
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Sequence 7, Appli
Sequence 4, Appli
Sequence 5, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 607, App
Sequence 607, App
Sequence 2790, App
Sequence 429, App
Sequence 4, Appli
Sequence 277, Ap
Sequence 277, Ap
Sequence 3, Appli
Sequence 37, Appli
Sequence 277, Ap
Sequence 277, Ap
Sequence 379, App
Sequence 2854, Ap
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Patent No. US20020132976A1

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RESULT US-09-		ი	C	C			O		a	C			O								o	C				
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		Sequence 31, Appl Sequence 73, Appl	29,	Sequence 14, Appi	1047, /	1235,	699,						Sequence 1, Appli								335,		257,	Sequence 11130, A	Sequence 12564, A	

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RESULT 2
US-09-755-100-4
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Sequence 4, Application US/09755100
Patent No. US2002009189A1
GENERAL INFORMATION:
APPLICANT: SAUTIZKY, Kinneret et al.
TITLE OF INVENTION: NOVEL NUCLEIC ACID
FILE REFERENCE: 2786-0156P
                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 7
LENGTH: 24
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Chong, Pele
APPLICANT: Pedyzzak, Artur
APPLICANT: Sia, Charles Dwo Yuan
TITLE OF INVENTION: Immunogenic Peptides Derived
TITLE OF INVENTION: Uses Thereof
FILE REFERENCE: 11014-24
                                                                                                                                                                                                                                                        Query Match 100
Best Local Similarity 100
Matches 24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 60/195,456
PRIOR FILING DATE: 2000-04-10
NUMBER OF SEQ ID NOS: 12
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/829,004A CURRENT FILING DATE: 2001-04-16
                                                                                                                                                                                 100.0%; Score
100.0%; Pred.
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Mismatches
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                      AND AMINO ACID SEQUENCES
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Sequence 5, Application US/09755100

Patent No. US20020099189A1

GENERAL INFORMATION: SEQUENCES AT TITLE OF INVERTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES FILE REFERENCE: 2786-01569

CURRENT APPLICATION NUMBER: US/09/755,100

CURRENT FILING DATE: 2001-01-08

PRIOR APPLICATION NUMBER: IL 128587

PRIOR APPLICATION NUMBER: IL 128587

PRIOR APPLICATION NUMBER: IL 128587

PRIOR FILING DATE: 1999-02-18

PRIOR APPLICATION NUMBER: IL 131363

PRIOR APPLICATION NUMBER: IL 131363

PRIOR FILING DATE: 1999-08-11

NUMBER OF SEQ ID NOS: 12

SOFTWARE: PATENTIN Ver. 2.1

SEGO ID NO 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/755,100
CURRENT FILING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: IL 128587
PRIOR FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: IL 129439
PRIOR FILING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: IL 131363
PRIOR APPLICATION NUMBER: IL 131363
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 12
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO. 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-755-100-5
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                                                                                                                                                                                                             Sequence 3, Application US/09755100 Patent No. US20020099189A1 GENERAL INFORMATION:
              APPLICANT: SAVITZKY, Kinneret et al.
TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REFERENCE: 2786-0156p
CURRENT APPLICATION NUMBER: US/09/755,100
CURRENT FILING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: IL 128587
PRIOR APPLICATION NUMBER: IL 129439
PRIOR FILING DATE: 1999-04-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 100.
Best Local Similarity 100.
Matches 24; Conservative
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ORGANISM: Homo sapiens
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APPLICATION NUMBER:
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24; Conservative
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100.0%; Pred. No.
tive 0; Mismatch
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100.0%; Pred. No. 0.026;
  IL 131363
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; TYPE: DNA; ORGANISM: Homo sapiens US-09-755-100-1
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; ORGANISM: Homo sapiens
US-09-755-100-3
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US-09-755-100-2
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US-09-755-100-1
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Patent No. US20020099189A1
GENERAL INFORMATION:
APPLICANT: SAVITARY, Kinneret et al.
TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REFERENCE: 2786-0156P
CURRENT APPLICATION UNMBER: US/09/755,100
CURRENT FILING DATE: 2001-01-08
                                                                                                                                                                                                                                                                                                                     Sequence 2, Application US/09755100 Patent No. US20020099189A1 GENERAL INFORMATION:
            PRIOR APPLICATION NUMBER: IL 131363
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 12
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
LECTION 1661
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 12
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
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Best Local Similarity 100.0%;
Matches 24; Conservative 0
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NUMBER OF SEQ ID NOS: 12
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 3
LENGTH: 3846
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Best Local Similarity 100.
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PRIOR TILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: IL 129439
PRIOR ETLING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: IL 131363
PRIOR ETLING DATE: 1999-08-11
PRIOR ETLING DATE: 1999-08-11
                                                                                                                           CURRENT FILING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: IL 128587
PRIOR FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: IL 129439
PRIOR FILING DATE: 1999-04-14
                                                                                                                                                                                                                  APPLICANT: SAVITZKY, Kinneret et al.
TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REFERENCE: 2786-0156P
CURRENT APPLICATION NUMBER: US/09/755,100
CURRENT FILING DATE: 2001-01-08
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os: 12
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Pred. No. 0.029;
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ORGANISM: Homo sapiens

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RESULT 8
US-09-780-669-607
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US-09-759-143-607
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                                                                                                                                                   Sequence 607,
Patent No. US
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SEQ ID NO 607
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Patent No. US20020022248A1
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                                         APPLICANT:
                                                                   APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C23
CURRENT FILING DATE: 2001-01-12
CURRENT FILING DATE: 2001-01-12
NUMBER OF SEQ ID NOS: 934
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TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                   07, Application US/09780669
US20020051977A1
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Fanger, Gary R. Retter, Marc W. Stolk, John A.
                                                                              Dillon, Davin C.
Mitcham, Jennifer L.
Harlocker, Susan L.
                                         Henderson, Robert A. Kalos, Michael D.
                                                                  Jiang, Yuqu
                                                                                                                         Xu, Jiangchun
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100.0%; Pred. No. 0.0
tive 0; Mismatches
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Pred. No. 1.1;
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RESULT 10
US-09-974-300-2790/c
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; OTHER INFORMATION: Primer
US-09-822-827-607
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US-09-822-827-607
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          Sequence 2790, Application US/09974300
Patent No. US20020146721A1
GENERAL INFORMATION:
APPLICANT: Berka, Randy M.
APPLICANT: Clausen, Ib Groth
TITLE OF INVENTION: Methods For Monitoring Multiple Gene
TITLE OF INVENTION: Expression
                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.534C1
CURRENT APPLICATION NUMBER: US/09/822.827
CURRENT FILING DATE: 2001-03-28
NUMBER OF SEQ ID NOS: 982
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 607
LENGTH: 22
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Best Local Similarity
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/780,669
CURRENT FILING DATE: 2001-02-09
NUMBER OF SEQ ID NOS: 943
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APPLICANT: Houghton, Raymond L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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OTHER INFORMATION: Primer
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Hepler, William
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Carter, Darrick
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ION: Expression 10085.500-US
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Pred. No.
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CURRENT APPLICATION NUMBER: US/09/974.300
CURRENT FILING DATE: 2001-10-05
PRIOR APPLICATION UNMBER: 09/680.598
PRIOR FILING DATE: 2000-10-06
PRIOR APPLICATION UNMBER: 60/279,526
PRIOR FILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 8481
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2790
LENGTH: 612
TYPE: DNA
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US-09-764-868-429/c
Sequence 429, Application US/09764868
Patent NO. US20020168711A1
GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PTZ32
                                                              RESULT 12
US-09-036-613-4/c
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Best Local Similarity
"~+~hes 18; Conserv?
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; ORGANISM: Bacillus licheniformis
US-09-974-300-2790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/764,868
CURRENT FILING DATE: 2001-01-17
Prior application data removed - refer to PALM or
NUMBER OF SEQ ID NOS: 1510
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 429
LENGTH: 1484
           Sequence 4, Application US/09036613 Patent No. US20020115135A1 GENERAL INFORMATION:
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Best Local Similarity 87.5%;
Matches 21; Conservative
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LOCATION: (10)
OTHER INFORMATION: n equals a,t,g, or c
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OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: SITE
LOCATION: (9)
OTHER INFORMATION: n equals a,t,g, or c
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                                                                                                                                                                                                                                                                                           OTHER INFORMATION: n equals a,t,g, or c
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APPLICANT:
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Pred. No. 3
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RESULT 13
US-09-664-761-16219
Sequence 16219, Application US/09864761
Sequence 16219, Application US/09864761
Sequence 16219, Application US/09864761
Patent NO. US20020048763A1
FARENT INFORMATION:
APPLICANT: Hank, David K.
APPLICANT: Hankel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REFERENCE: Acomica-x-1
CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
CURRENT FILING DATE: 2001-05-23
CURRENT FILING DATE: 2001-05-23
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PILING DATE: Filed Herewith
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
APPLICATION NUMBER:
FILING DATE:
APPLICATION NUMBER:
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: PF-04
REFERENCE/DOCKET NUMBER: PF-04

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PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR FILING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: US 024263.6
PRIOR APPLICATION NUMBER: US 60/236,359
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
OPERATING SYSTEM: DOS
SOFTWARE: FastsP0 for Windows Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
IMMEDIATE SOURCE:
LIBRARY: BRAINOS12
CLONE: 2762136
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APPLICANT: Corley, Neil C.
TITLE OF INVENTION: SYNAPS
NUMBER OF SEQUENCES: 7
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CITY: Palo Alto
STATE: CA
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FILING DATE:

2001-01-30

APPLICATION NUMBER: PCT/US01/00667 APPLICATION NUMBER: PCT/US01/00664

APPLICATION NUMBER: PCT/US01/00666

2001-01-30 2001-01-30

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APPLICANT: Harper, Jeff
APPLICANT: Kreps, Joel
APPLICANT: Wang, Xun
APPLICANT: Zhu, Tong
APPLICANTON: SAME, AND METHODS OF USE
FILE REFERENCE: SCRIP1300-3
CURRENT APPLICATION NUMBER: US/09/938,842A
CURRENT FILING DATE: 2001-08-24
PRIOR APPLICATION NUMBER: US 60/227,866
PRIOR APPLICANTON NUMBER: US 60/264,647
PRIOR APPLICANTON NUMBER: US 60/264,647
PRIOR APPLICATION NUMBER: US 60/300,111
PRIOR FILING DATE: 2001-06-22
NUMBER DE SEO. TO NOS. 3770
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; Sequence 2277, Application US/09938842A

; Patent No. US20020160378A1
; TYPE: DNA
; ORGANISM: Arabidopsis thaliana
US-09-938-842A-2277
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; OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL US-09-864-761-16219
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LENGTH: 569
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                                                                                                                      PRIOR FILING DATE: 200 NUMBER OF SEQ ID NOS:
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OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00663
OR FILING DATE: 2001-01-30
OR APPLICATION NUMBER: PCT/US01/00662
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OR APPLICATION NUMBER: PCT/US01/00661
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FILING DATE: 2000-09-21
APPLICATION NUMBER: US 09/608,408
FILING DATE: 2000-06-30
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Search completed: November 19, 2002, 10:16:24 Job time : 90.2308 secs
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CURRENT FILING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: I TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                             SOFTWARE: FastSEQ
                                                          27025 TGTGGGTCCCAGGTGCCTTCC
                                                                                                                                                                                                        NAME/KEY: misc_feature LOCATION: (1)...(33239)
                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Human
                                                                                                                                                                                          OTHER INFORMATION: n =
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RESULT 1 AX281251 LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM REFERENCE AUTHORS TITLE JOURNAL	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	Result No. Sc
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                                                                                                                                                                                                                                                                                                          Submitsed (09-JAN-1989) Digby M.R., The Howard I Submitted (09-JAN-1989) Digby M.R., The Howard I Melbourne University, Parkville, Victoria 3052, 2 (bases 1 to 287)
2 (bases 1 to 287)
Digby, M., Zhang X.Y. and Richards, R.I.
Human prostate specific antigen (PSA) gene: struct the kalikrein-like gene, NGK-1
Nucleic Acids Res. 17 (5), 2137 (1989)
89183632
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Sequence 1 from patent US
AR047835
AR047835.1 GI:5969300
                                                                                                                                                                                                       Human PSA gene for
X13941
X13941.1 GI:35724
kallikrein; prosta
                                                                                                                                                                                                                                                                                                                                                       27; Conserv
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Location/Qualifiers
1. .200
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1 (bases 1 to 200)
Gundling.G.J.
Rapid RNA isolation procedure in the
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 287)
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kindly reviewed (16-may-1989) by Digby M.R. Location/Qualifiers
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PSA gene for prostate specific antigen
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Location/Qualifiers
1. .27
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58 c 67 g
                                                                                                                                                                                                       prostate specific antigen; PSA gene
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Pred. No. 0.048;
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9
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218 GTTCTGGTGCACCCCCAGTGGGTCCTC 244
                 1 GTTCTGGTGCACCCCCAGTGGGTCCTC 27
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27; Conservative
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288. ->287
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100 c 78 g 56 t
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ESULT 4
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FINITION PSA-prostate-specific antigen [human, breast cancer specimen, mRNA ccession 575755
ERSION 575755
ERSION 575755
ERSION 575751 GI:861469
EXWORDS

ON S75755.1 GI:861469

DRDS Homo sapiens breast cancer specimen.

E Homo sapiens breast cancer specimen.

Eukaryota; Metazoa; Chordata; Crania;
Mammalia; Eutheria; Primates; Catarri

RENCE 1 (bases 1 to 569)

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

E 1 (bases 1 to 569)

S Monne, M., Croce, C.M., Yu, H. and Diamandis, E.P.
Molecular characterization of prostate-specific antigen messenger RNA expressed in breast tumors

RNA expressed in breast tumors

L Cancer Res. 54 (24), 6344-6347 (1994)

RNA expressed in breast tumors Cancer Res. 54 (24), 6344-6347 (1994) 95079406 7527295 GenBank staff at the National Library of Medicine created this entry (NCBI gibbsq 161505) from the original journal article.

Location/Qualifiers
1.569
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Homo sapiens mRNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 658)
Riegman, P. H., Klaassen, P., van der Korput, J.A., Romijn, J.C. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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glycoprotein; prostate antigen; prostate-specific antigen;
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708 bp for putative p
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(PA) gene,
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clone PA 424.
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Sequence 4
AR153342
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194 GTTCTGGTGCACCCCCAGTGGGTCCTC 220
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Characterization of PSA-RP2, a protein related to prostate-specific antigen and encoded by alternative.hKLK3 transcripts
Dir. J. Biochem. 268 (16), 4408-4413 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AJ310937
AJ310937.1 GI:14422304
alternative splicing; HKLK3 gene; kallikrein 3; preproPSA-RP2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              related accession number AJ310938.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (bases 1 to 708)
                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                           /gene="KLK3"
/evidence=experimental
/ 214 c 219 g
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<1. .37
                                                                                                                                                                                                                                                                                                                                                                                                                                                               /gene="KLK3"
708
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578. .708
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110. .577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /product="putative prepropsa-rp2"
/protein_id="CAC41631.1"
/protein_id="CAC41631.1"
/db_xref="Gi:14422305"
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MSLLKNRFLRSPGDDSSHDLMLLRLSEPAELTDAVKVMDLPTQEPALGTTCYASGWGSI
MSLLKNRFLRSPGDDSSHDLMLLRLSEPAELTDAVKVMDLPTQEPALGTTCYASGWGSI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /gene="KLK3"
/function="unknown"
/note="alternative splicing, transcript
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:9606"
/clone="NH101007.97"
/cell_line="LNCap"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene="KLK3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /codon_start=1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /gene="KLK3"
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711 bp
from patent US 6235486.
                                                                                                                                                                                                                                                                                 100.0%; Score 27; DB 9; 100.0%; Pred. No. 0.049;
                                                                                                                                                                                                                                                          0;
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                                                                                                                                                                                                                                                                                                            Length 708
                               linear
                               PAT 08-AUG-2001
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RESULT 9
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KEYWORDS
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Best Local S
Matches 27
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E 1 (bases 1 to 711)

S Tindall, D. J., Young, C.Y.F., Mccormic, D. J., Klee, G.G., Saedi, M Kumar, A., Rittenhouse, H.G. and Wolfert, R.L.

Method for detecting metastatic prostatic cancer

L Patent: JP 2001503991-A 14 27-MAR 2001;

MAYO FOUNDATION FOR MEDICAL EDUCATION AND RESEARCH, HYBRITECH:

OS Unidentified

PN JP 2001503991-A/14

PD 27-MAR-2001

PP 14-NOV-1997 JP 1998522888

PR 14-NOV-1997 JP 1998522888

PR 14-NOV-1997 JF 1998522888
                                                   GTTCTGGTGCACCCCCAGTGGGTCCTC 27
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PC C1
G01N33,
PC G0
CC St
CC TC
FH K6
FT SC
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Location/Qualifiers
                                                                                                     ch 100.0%;
l Similarity 100.0%;
27; Conservative 0
                                                                                                                                                                                                                                                                                                                                                           GEORGE
PI MC
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JP 2001503991-A/14.
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                                                                                                                                                                                                                                                                               Topology: Linear;
                                                                                                                                                                                                                                                                                           Strandedness: Double;
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L WOLFERT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    for detecting metastatic
                                                                                                                                                                                                           /organism-'Unidentified'.
Location/Qualifiers
1. 711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="unknown"
208 c 206 g
                                                                                                                                                                      /organism="unidentified"
/db_xref="taxon:32644"
208 c 206 g 14
                                                                                                        0;
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                                                                                                     Score 27; DE
Pred. No. 0.0
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA linear prostatic cancer.
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0.049;
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0.049;
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                                                                                                        Gaps
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VERSION
KEYWORDS
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VERSION
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HSA459782
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27; Conserv
                                                                                                         Homo sapiens variant 1.
                                       Homo sapiens
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             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                  alternative splicing; KLK3 gene;
                                                                              AJ459782.1
                                                                                                                                   HSA459782
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                                                                                                                                                                                                                                                                                                                     185
                                                                              GI:20520640
                                                                                                                     mRNA
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Courty,Y.
Direct Submission
Submitted (03-MAY-2002) Courty Y., EMI-U 0010, Laboratoire
Submitted (03-MAY-2002) Courty Y., EMI-U 0010, Laboratoire
d'Enzymologie, Universite F. Rabelais, 2 bis bvd Tonnelle,
37032 cedex, FRANCE
37032 cedex, FRANCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Heuze-Vourc'h,N. and Courty,Y. Complex alternative splicing of the hKLK3 gene coding for the tumour marker PSA (prostate-specific-antigen) unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AJ459783.1 GI:20520642 alternative splicing; KLK3 gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens mRNA for prostate
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Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (bases 1 to 870)
                                                                                 100.0%;
ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                                            /gene="KI
94..804
                                                                                                                                                                                                                                                                                                /Translation="MWVPVVFITLSVTWIGAAPLILSRIVGGWECEKHSQPWQVLVAS
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EPEEFLTPKKLQCVDLHVLSNDVCAQVHPQKVTKFMLCAGRWTGGKSTCSGDSGGPLV
CQGVLQGITSWGSEPCALPERPSLYTKVVHYRKWIKDTIVANP"
                                                                                                                                                                            /product="prostate specific antigen" 257 c 242 g 186 t
                                                                                                                                                                                                                                                                                                                                                                                                 /product="prostate specific antigen"
/protein_id="CAD30845.1"
/db_xref="GI:20520643"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="YC140405-00"
                                                                                                                                                                                                                         'gene≖"KLK3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                            /evidence=experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 codon_start=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'note="alternative splice variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gene="KLK3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /gene="KLK3"
22. .807
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /tissue_type="prostate"
L. .870
                                                                                                                                                                                                                                                              "KLK3"
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for

876 bp

mRNA specific

linear antigen (

(KLK3 gene), splice

prostate

specific antigen

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                                              Direct Submission
Submitted (08-NOV-1994) Jose M J G Moreno,
Jefferson University Hospital, 1025 Walnut
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Submitted (03-MAY-2002) Courty Y., EMI-U 0010, Laboratoire
d'Enzymologie, Universite F. Rabelais, 2 bis bvd Tonnelle,
                                                                                                                                               2 (bases 1 to 990)
Moreno, J.M.
                                                                                                                                                                                                     A Comparative Analysis of Prostate Specific Antigen Gene Sequence in Benign and Malignant Prostate Tissue Unpublished
                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 990)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human prostate specific antigen precursor mRNA, complete cds \tt u17040
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                                                                                                                                                                                                                                                                                                  Gomella,L.G.
                                                                                                                                                                                                                                                                                                                               Monne, M.M., Moreno, J.M., Mele, C.M.; Mulholland, G.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Courty, Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Unpublished
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Complex alternative splicing of the hKLK3 gene coding
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/gene="KLK3".
94. .681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /product="prostate specific antigen"
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/db_xref=="Ga1:20520641"
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/translation="MWVPVVFLT"
/translation="MWVPVVFLT"
/translation="MWVPVVFLT"
/translation="MWVPVVFLT"
/translation="MWVPV
Location/Qualifiers
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235 c 231 g 190 t
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22. .684
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/db_xref="taxon:9606"
/clone="YC171105-00"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /gene="KLK3"
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0.049;
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Street, P
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                                                        Philadelphia, PA
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Croce,C., Gomella,L., Mulholland,S.Grant.,
Fischer,R.
                                                                                                                                                                                                                                                                                                                                                                                 992 bp I
Sequence 13 from patent US 5674682.
167863
167863.1 GI:2829985
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                                                                                                                                                                                                                                                                                                                                                                 Unknown
                                                                                                                                                                                                                                                                       Patent:
                                                                                                                                                                                                                                                                                    cancer
                                                                                                                                                                                                                                                                                             Nucleic acid primers for detecting micrometastasis of prostate
                                                                                                                                                                                                                                                                                                                                                        Unknown
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114. .8
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EPEEFLTPKKLQCVDLHVISNDVCAQVHPQKVTKFMLCAGRWTGGKSTCSGDSGGPLV
CNGVLQGITSWGSEPCALPERPSLYTKVVHYRKWIKDTIVANP"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /product="prostate
93. .113
                                                                                                                                                                                                                       /organism="unknown"
291 c 272 g
                                                                                                                                                                                                                                                         Location/Qualifiers
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291 c 273 g
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RGRAVCGGVLVHPQWVLTAAHCIRNKSVILLGRHSLFHPEDTGQVFQVSHSFPHPLYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /codon_start=1
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12. .827
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                                             linear PRI 13-MAY-
ant 1 mRNA, complete
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
Homo sapiens
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 1130)
David, A., Mabjeesh, N., Azar, I., Biton, S., Engel, S., Bernstein, J.,
Romano, J., Avidor, Y., Waks, T., Eshhar, Z., Langer, S.Z.,
Lifschitz-Mercer, B., Matzkin, H., Rotman, G., Toporik, A., Savitsky, K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ch 100.0%; Score 27; DB 9; Length 1130;
1 Similarity 100.0%; Pred. No. 0.59; Length 1130;
27; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                          Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 1350) Mitcham, J.L., Harlocker, S.L., Jiang, Y., Xu, J., Dillon, D.C., Mitcham, J.L., Harlocker, S.L., Jiang, Y., Redo, S.G., Kalos, M.D., Fanger, G.R., Day, C.H., Retter, M.W., Stolk, J.A., Skelky, Y.A., Wang, A. and Meagher, M.J.
                                                                                                                                                                                                                                                                                                                                                                                       AX200986 1 GI:15390813
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Submitted (11-JAN-2001) Compugen Ltd.,
69512, Israel
                                                                                                                        Patent: WO 0151633-A 616 19-JUL-2001; CORIXA CORPORATION (US)
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vid,A., Engel,S., Azar,I., Bernstein,J.,
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                  284
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/codon_start=1
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                                                                                Location/Qualifiers
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404 c 391 g 27
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/map="19q13.3"
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AUTHORS
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Sequence 616 from Patent W00173032.
AX267642
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Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                     Patent: WO 0173032-A 616 04-OCT-2001; CORIXA CORPORATION (US)
                                                                                                                                                                                                                                                                                                                               Xu,J., Dillon,D.C., Mitcham,J.L., Harlocker,S.L., Jiang,Y., Kalos,M.D., Fanger,G.R., Retter,M.W., Stolk,J.A., Day,C.H., Vedvick,T.S., Carter,D., Li,S.X., Wang,A., Skelky,Y.A., Hepler,W.T. and Henderson,R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AX267642.1 GI:16516315
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Copyright (c) 1993 - 2002 Compugen Ltd
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## ALIGNMENTS

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                                              (AVET ) AVENTIS PASTEUR LTD
                                                                                                                                               10-APR-2000; 2000US-195456P
                                                                                                                                                                                                                                                 10-APR-2001; 2001WO-CA00473
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The present invention describes a method for detecting the occurrence of a nucleic acid synthesising reaction during the synthesis of a nucleic acid with an enzyme. The method comprises the use of an insoluble substance formed in the reaction as an indication. The method can be used for detecting the presence or absence of a nucleic acid synthesising reaction (especially a nucleic acid amplification reaction). The method is useful in gene analysis, diagnosis of cancer and other diseases, and
                                                                                                                                                                                                                                                                                                   Detecting the product of an enzyme-based nucleic acid synthesizing reaction, useful in gene analysis, diagnosis of cancer and other diseases, and in identifying bacteria, comprises the formation of an insoluble substance -
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AAV60379

AAV60379 standard; mRNA;

ВP

В Q

39

GTTCTGGTGCACCCCCAGTGGGTCCTC

27 65

GTTCTGGTGCACCCCCAGTGGGTCCTC

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Best Local S
Matches 27
 Matches
                       Query Match
                                                                    reaction. The method uses an arbitrary primer to initiate the complementary chain synthesis reaction. The method is particularly useful in gene and disease diagnosis. It is a highly efficient and reaction specific method in which no temperature variation is required. The present sequence is used in an example illustrating the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                in identifying bacteria. The present sequence oligonucleotide used in the exemplification of
                                                                                                                                   The invention relates to a method for synthesising a nucleic acid using a double-stranded nucleic acid as template and incubating under conditions allowing the establishment of a complementary chain synthesis
                                                                                                                                                                                                      Isothermal amplification of nucleic acids using double-stranded nucleic acid as template to establish complementary chain synthesis reaction from primer enabling base pairing in domain to be annealed, useful e.g. in gene diagnosis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 178 BP; 33 A; 56 C; 50 G;
                                              Sequence 178 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABA01120;
                                                                                                                                                                                Example 1; Page 63; 75pp; Japanese.
                                                                                                                                                                                                                                                                   WPI; 2002-010907/01.
                                                                                                                                                                                                                                                                                          Notomi T,
                                                                                                                                                                                                                                                                                                                                        07-APR-2000; 2000JP-0111939
                                                                                                                                                                                                                                                                                                                                                               30-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABA01120
                                                                                                                                                                                                                                                                                                                (EIKE ) EIKEN KAGAKU KK
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            Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PSA nucleotide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nucleic acid synthesis; complementary chain synthesis; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 Similarity
27; Conserv
            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard;
                                                                                                                                                                                                                                                                                          Nagamine
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 Conservative
                                                                                                                                                                                                                                                                                                                                                               2001WO-JP02771
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                                                33 A; 56 C; 50 G; 39 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
           100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      178
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 0,
           Score 27;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 27; D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        39 T; 0 other;
                                                                                                                                                                                               ...
            DB 24;
0.034;
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0.034;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of the presents
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                        Length 178;
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Gaps
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PF XXX
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                                                                                                                                                                                                                                                                                                                                 Best
                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents a mRNA target sequence from exons 2 and 3 of the prostate specific antigen (PSA) gene... The sequence is used to exemplify the invention. The specification describes a method for purifying RNA. The method comprises contacting a test sample with polyvalent transition metal ions to form a precipitant and a supernatant, separating the precipitant from the supernatant, and collecting the supernatant to obtain a purified solution of total RNA. The RNA is used for nucleic acid amplification and disease detection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Purification of RNA, used for nucleic acid amplification and disease detection - by precipitating other sample components with transition metal ions, separating and collecting the purified RNA
                                                                                         Diagnosis; prostate cancer; prostate specific antigen; PSA; metastatic prostate cancer; prostate cell; secondary tumour; bone metastatic anchoring; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 5; Columns 9-10; 6pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gundling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mRNA
16-APR-1997;
                                                                    Homo
                                                                                                                                                                                                             AAT92975 standard; cDNA; 208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US5817798-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ^{\text{ds}}
                       23-OCT-1997
                                             WO9739139-A1
                                                                                                                                        cDNA for prostate
                                                                                                                                                                                       AAT92975
                                                                                                                                                                                                                                                                                                                                                                  Sequence 200 BP; 31 A; 58 C; 67 G; 44 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-556473/47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-OCT-1998
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                                                                                                                                                                 24 - APR - 1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo
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                                                                                                                                                                                                                                                                                                                                 Local
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                                                                   sapiens
                                                                                                                                                                                                                                                                     GTTCTGGTGCACCCCCAGTGGGTCCTC 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        target sequence; prostate specific antigen; PSA; RNA purification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               target
                                                                                                                                                                                                                                                                                                                    l Similarity
27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence from exons 2 and 3 of PSA gene
                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LAB.
97WO-US06497.
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                                                                                                                                         specific
                                                                                                                                                                                                                                                                                                                                 100.0%;
                                                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                         antigen
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                                                                                                                                                                                                                                                                                                                                 Score 27;
Pred. No.
                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                         (PSA).
                                                                                                                                                                                                                                                                                                                                 0.035;
                                                                                                                                                                                                                                                                                                                                                                  0 other;
                                                                                                                                                                                                                                                                                                                                          DB 19;
                                                                                                                                                                                                                                                                                                                    0;
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                                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                   Gaps
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RESULT 6
AAA61482
Db
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 metastatic prostate cancer. Detection of PSA mRNA in the bloodstream indicates that prostate cells are circulating in the blood and confirms diagnosis of cancer. Circulating prostate cells also indicate a risk of a secondary tumour or bone metastatic anchoring. Detection of the PSA mRNA after prostatectomy or radiotherapy indicates a risk that the prostate cancer has spread and that the surgery and radiotherapy was not effective. PCR primers AAT92971-73 were used for the monitoring and diagnosis of patients with prostate cancer. The method comprises the detection of PSA mRNA by reverse transcriptase polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Monitoring or diagnosis of prostate cancer - by detecting prostate-specific antigen mRNA using specific primers
                                                                                                                                                                                                                                                                                                                                             AAA61482 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDNA sequences AAT92974-77 er
The detection of PSA mRNA in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure;
                                                                                                                                                                                                                                                                               Prostate
                                                                                                                                                                                                                                                                                                   23-OCT-2000
                                                                                                                                                                                                                                                                                                                        AAA61482;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1997-526473/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SMIK ) SMITHKLINE BEECHAM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                   133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (RT-PCR)
                                                                                                                                                                                                                                                                                                                                                                                                                       GTTCTGGTGCACCCCCAGTGGGTCCTC
                                                                                                                                                                                                                                                                                                                                                                                                  GTTCTGGTGCACCCCCAGTGGGTCCTC 159
                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
27; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      208
                                                                                                                                                                                                                                                                              specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B₽;
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                                                                                                                                                                                                                                                                              antigen gene fragment.
                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; 62 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25pp;
                                                                                                                                                                                                                                                                                                                                              517
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  encode the prostate specific antigen (PSA) in peripheral blood is associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   English.
                                                                                                                                                                                                                                                                                                                                              ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     73 G; 44 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 27;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                       27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.035;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0 other;
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Homo sapiens
                                                     Isothermal DNA amplification; nucleic acid synthesis; PSA gene template; prostate specific antigen; human; c
                                                           ds
                                                                                     detection;
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Nucleic acid synthesis using primer containing a sequence identical to one on the template for isothermal amplification using simple reagents

WPI; 2000-376587/32

Notomi

Ŧ,

Hase T;

09-NOV-1998; 08-NOV-1999;

98JP-0317476 99WO-JP06213

(EIKE ) EIKEN KAGAKU KK

18-MAY-2000

Example

8;

Fig 17;

95pp; Japanese

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RESULT 7
ADAHA1045
ID AAHA
AC AAHA
AC AAHA
AX 30-AA
XX Pros
XX Pros
XX WO2C
XX Unid
PN WO2C
XX Unid
PN WO2C
XX Unid
PN WO2C
XX UNId
PF 28-P
XX O8-P
XX O8-P
XX UNId
XX UNId
PT Synt
XX Exar
XX
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CA region is produced at the 3' end of a target sequence which contains sub-sequences designated F1, F2C and F1C, where F1C is the complement of F1. A loop is formed by annealing of F1 to its complementary sequence F1C, and a primer containing a sub-sequence F2 (complementary strand to the template. The complementary strand to the template is synthesised using a DNA polymerase, and the process is repeated using sub-sequences at the 3' end of the complementary strand designated R1, R2C and R1C, using a primer containing R2 at its 3' end. Cand methods for the detection of a nucleic acid sequence in a sample cusing a labeled probe containing a sequence complementary to the loop cregion of the target sequence. The method of the invention provides an efficient and highly specific method for isothermal amplification of a target nucleic acid sequence using a simple reagent mixture.

CC of a target nucleic acid sequence using a simple reagent mixture.

CC of starget nucleic acid sequence using a simple reagent mixture.

CC of a target nucleic acid sequence using a simple reagent mixture.

CC of a target nucleic acid sequence using a simple reagent mixture.

CC of a target nucleic acid sequence using a simple reagent mixture.
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Best Local
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The resent invention describes a new method for synthesising nucleic acids comprising: (a) preparing a nucleic acid, with an F2c region, able to form a loop by annealing the F1 region of the 3' end to the F1c region on the same chain; (b) synthesis of a complementary chain using the 3' end of F1 as the starting point; (c) replacing the complementary to F2c) at its 3' end and using it as a starting point for complementary chain synthesis by a polymerase; and (d) replacing the chain synthesised in step c by annealing a polynucleotide containing a region complementary to a part of the chain and using its 3' end as the starting point. Also described are: (1) a kit for synthesising nucleic acids; and (2) a kit for detecting target base sequences. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 8; Fig 17; 96pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthesizing nucleic acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Notomi T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (EIKE ) EIKEN KAGAKU KK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prostate specific antigen; nucleic acid synthesis; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Prostate specific antigen related nucleotide sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GITCTGGTGCACCCCCAGTGGGTCCTC 207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2001-343603/36.
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27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         517 BP; 94 A; 163 C; 151 G; 109 T; 0 other;
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Pred. No.
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0.038;
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RESULT 8
AAX08947
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Best Local S
Matches 27
                                    Human kallikrein 2 (hK2) is expressed at elevated levels relative to the prostate cancer antigen, prostate specific antigen (PSA) by breast cancer cells when stimulated by androgens. Detecting levels of hK2 may provide a simple diagnostic tool for detecting or determining breast cancer. Detecting hK2 is achieved by producing an hK2 DNA from hK2 RNA by reverse transcription. The hK2 DNA obtained is then amplified by PCR and detected using probes. Determination of breast cancer may also be determined by exposing the hK2 DNA by periode to an agent which binds to it and then detecting the binary complex formed. The amount of complex formed correlates directly to the presence of breast cancer in the human individual. The methods can be used in the diagnosis, treatment and/or monitoring of the progression or remission of breast cancer and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence represents a prostate specific antigen related nucleotide sequence, which is used in an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human kallikrein 2; hK2; breast cancer; diagnosis; detection; treatment; monitoring; prostate specific antigen; PSA; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 517 BP; 94
                           monitoring of the progression monitoring hK2 levels.
                                                                                                                                                                                                                                                       Disclosure; Page 34-35;
                                                                                                                                                                                                                                                                                    Use of human kallikrein 2 - as a marker for developing products for the diagnosis, prognosis, monitoring and treatment of breast cancer
                                                                                                                                                                                                                                                                                                                                                                                 Klee GG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-DEC-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX08947 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                            (KLEE/) KLEE G G.
(MAYO-) MAYO FOUNDATION.
(TIND/) TINDALL D J.
(YOUN/) YOUNG C Y F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human prostate specific antigen (PSA) gene.
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l Similarity 100.0%;
27; Conservative 0
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                                                                                                                                                                                                                                                       67pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
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Pred. No. 0.038;
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Sequence 711 BP; 148 A;

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206 <u>و</u>

149 T; 0

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RESULT 9
ABK86205
The invention relates to a method of inducing a cell-mediated immune cresponse against a cell comprising a target antigen (I) in a subject, treating a subject having undesired cells, for example tumour cells or virally infected cells (C), reducing the number of (C) in a subject, and stimulating presentation of (I) by a cell. This is done by administering a polynucleotide (II) encoding a variant of (I), so that (II) expressed in a cell and cell-mediated immune response is induced. The method can be used to treat prostate cancer, breast cancer, colorectal cancer and pancreatic cancer, as well as lymphomas and leukaemias. The method is also useful in treating chronic viral infections such as those caused by hepadnaviruses, lentiviruses (including human immunodeficiency virus (HIV)), herpesviruses and the flaviviruses and pestiviruses. The present sequence represents the coding sequence of human prostate specific antigen (PSA) variant which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local
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                                                                                                                                                                                                             Disclosure; Page 121-122; 146pp; English.
                                                                                                                                                                                                                                                      a cell,
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                                                                                                                                                                                                                                                                                                                                                  Mincheff MS,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABK86205 standard; cDNA; 744
                                                                                                                                                                                                                                                                   reducing
                                                                                                                                                                                                                                                                                 Inducing
                                                                                                                                                                                                                                                                                                                                                                                                                     (AMBI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunostimulant;
virus infection;
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                                                                                                                                                                                                                                                                                                                                                                                             [LOUK/)
                                                                                                                                                                                                                                                                                                                                                                                                        (MINC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; prostate specific antigen; PSA; cytostatic; antiviral; immunostimulant; cell-mediated immune response; tumour; breast crizes infection; prostate cancer; colorectal cancer; pancreatic (lymphoma; leukaemia; hepadnavirus; lentivirus; herpesvirus; human immunodeficiency virus; HIV; flavivirus; pestivirus; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  85
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DB; AAU98921.
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) ZOUBAK S.
                                                                                                                                                                                                                                                                                                                                                                                                      AMERICAN FOUND BIOLOGICAL MINCHEFF M S.
                                                                                                                                                                                                                                             g a cell-mediated immune response against a target
g undesired cells and stimulating presentation of a
comprises administering a polynucleotide encoding
yen -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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7..744
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Prostate specific antigen (PSA) variant with signal sequence deleted"
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                                                                                                                                                                                                                                                                                                                                                  Zoubak
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (PSA) variant
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g a variant
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RESULT 10
ABV25078
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                                                                                                                             a nucleotide sequence given in Tables 1-9 (ABV00010-ABV52213) or compensation or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate
                    patient;
(I) is a
                                                                                                                                                                                                                                                  Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, used for detecting presence of prostate cancer, stage of prostate cancer.
                                             cancer in a patient;
(d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient;
(e) selecting a composition for inhibiting prostate cancer in a patient (f) assessing the prostate cell carcinogenic potential of a compound;
(g) determining whether prostate cancer has metastasized in a patient;
(h) assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                       The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the
                                                                                                                                                                                                                             Claim 1;
                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                                                              Schlegel
                                                                                                                                                                                                                                                                                                                                                                                        09-JUN-2000;
18-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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25-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                                    (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             selecting a composition for inhibiting prostate cancer in a patient; assessing the prostate cell carcinogenic potential of a compound; determining whether prostate cancer has metastasized in a patient; assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the signal sequence deleted, used the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       prostate expression marker cDNA 25069
                        also
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27; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
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2000US-207454P.
2000US-211314P.
2000US-219007P.
2000US-255281P.
                        useful
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                       as a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytostatic; gene; ss.
                       pharmacodyanamic or
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                213 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                carcinogen;
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                       pharmacogenomic marker
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Query Match

Sequence

841

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262 C;

262 G;

173

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Score 27;

DB Τ;

23;

Length 841;

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RESULT 1:
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Matches
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Best Local :
                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                             The invention relates to a method of immunization against prostatic cancer (PC) by administering a nucleic acid (this sequence) that encodes prostate-specific antigen (PSA), or its immunologically active fragment, operably linked to regulatory sequences. The nucleic acid is taken up by cells and then expressed to generate an immune response against PSA. The method is used to protect against, or treat, PC, or more generally to induce a cellular immune response to PSA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAZ47134 standard; DNA; 990 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunization; prostate cancer; prostate-specific antigen; PSA;
cytostatic; immune response; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human prostate-specific antigen gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-MAR-2000 (first entry)
Prostate specific antigen; primer; human; prostate cancer;
                           Human prostate specific antigen gene.
                                                                                      AAT91054;
                                                                                                                  AAT91054 standard; DNA; 992 BP
                                                                                                                                                                                                                                                                                                         Sequence 990 BP; 209 A; 291 C; 273 G; 217 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 42; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunization against prostatic cancer using nucleic acid encoding prostate-specific antigen, for treatment or prophylaxis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-072551/06.
P-PSDB; AAY56048.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Weiner DB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYPE-) UNIV PENNSYLVANIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-MAY-1998;
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                                                                                                                                                                                                                                                 l Similarity
27; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kim JJ;
                                                                                                                                                                                                                                               100.0%; Score 27; DB 21; Length 990; Illarity 100.0%; Pred. No. 0.04; Conservative 0; Mismatches 0; Indels
                                                       (first entry)
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/*tag= u

/note= "PR8"

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550..569
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complement (1..19)
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231
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672..827
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1..87
                                                                                                                                               complement (685..702)
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/note= "PSA2"
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/note= "PR6"
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/note= "PR7"
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RESULT 13
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29-OCT-1992;
23-AUG-1994;
                        15-DEC-1994;
29-OCT-1992;
23-AUG-1994;
12-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antigen (PSA) cDNA PCR fragment. New oligonuclectide primers (AAT91044-53) are designed to amplify the PSA gene but not the human glandular kallikrein gene, which has 75-85% homology to the PSA gene. The primers can be used for diagnosis of prostate cancer micrometastasis. A claimed diagnostic kit for detecting prostate cancer comprises a pair of primers, where one of the primers is selected from those given in AAT91044-53, and a means for visualising amplified DNA (ethidium bromide stain, 32 por biotin). The invention also includes a method of detecting cells which express prostate cancer specific sequences by detecting the presence of nucleic acids specific for prostate cancer or a signal specific for
                                                                                                                                                           Homo
                                                                                                                                                                                Prostate specific antigen; PSA; D1 stage prostate cancer; D2 stage prostate cancer; D3 stage prostate cancer; human; diagnosis; glandular kallikrein gene; micrometastasis; detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This DNA sequence comprises a 992-bp human prostate specific antigen (PSA) cDNA PCR fragment. New oligonucleotide primer:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotide primers specific used for the diagnosis of prostat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-OCT-1997
                                                                                     12-DEC-1996;
                                                                                                            17-AUG-1999
                                                                                                                                  US5939258-A
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                                                                                                                                                                                                                                                                                   AAX99846;
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(UYJE-) UNIV
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                                                                                                                                                           sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                           (first entry)
 JEFFERSON THOMAS
                      94US-0358782.
92US-0973322.
94US-0294611.
96US-0764527.
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92US-0973322.
94US-0294611.
                                                                                     96US-0764527
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                                                                                                                                                                                                                                   antigen
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                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
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                                                                                                                                                                                                                                   PCR fragment.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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te cancer micro-metastasis
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Query Match
                                                                                                                                                    This sequence is a fragment of the human prostate specific antigen (PSA). The invention relates to a method of detecting prostate cancer micrometastasis in a patient having stage D1, D2, or D3 prostate cancer, comprising: (1) obtaining a sample nucleic acid from a patient tissue; (2) amplifying nucleic acids specific for prostate cancer or amplifying a signal generated by hybridisation of a probe specific to a prostate specific nucleic acid in the sample, comprising hybridisation to at least 1 prostate cancer specific perimer which is complementary to a prostate cancer specific gene which does not hybridise to the human glandular kallikrein gene; and (3) detecting the presence of the amplified nucleic acids or the amplified signal where the presence of the amplified nucleic acids or the amplified signal where the presence of the amplified nucleic
                                                                                                acids or signal indicates micrometastasis of prostate cancer. The meth is useful for detecting micrometastasis of prostate cancer. The method allows the detection of a low concentration of cells.
                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting micrometastasis of prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1999-468401/39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Croce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ç,
                                                         992 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fischer R,
                                                                                                                                                                                                                                                                                                                                                                                              Fig 2;
                                                           210 A; 291 C;
                                                                                                                                                                                                                                                                                                                                                                                              20pp;
100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gomella
                                                                                                                                                                                                                                                                                                                                                                                                English
Score 27;
Pred. No.
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                                                           272 G; 217 T; 2 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       cancer
                      DB
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                    20;
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                    Length
                                                                                                                                         The method
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RESULT 14
AAH99720
                                                                                                                                                   Matches
16-OCT-2001 (first entry)
                                               AAH99720 standard; cDNA; 1040
                                                                                                      200
                                                                                                                                                 Local Similarity
nes 27; Conserv
                                                                                                                            1 GTTCTGGTGCACCCCCAGTGGGTCCTC 27
                                                                                                      Conservative
                                                                                                                                                 0;
                                               ВP
                                                                                                                                                   Mismatches
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0

04;

0;

Human protein encoding cDNA sequence

SEQ ID

NO:555.

Ъ δÃ

23-DEC-1999; 21-JAN-2000; 25-APR-2000; antiinflammatory; antirheumatic; antiarthritic; immunosuppressive; antianterial; endocrine; cardiant; central nervous system; virucide; anti-HIV; fungicide; antimutagen; cardiovascular; antianaemic; anaemia; antiagregant; haemostatic; vulnerary; antiulcer; osteopathic; eczema; dermatological; antiallergic; antiasthmatic; antidabetic; cytostatic; neuroprotective; antidepressant; nootropic; antiparkinsonian; infection; immunostimulant; gene therapy; antisense therapy; vaccine; inflammation; antianaphylactic; rheumatoid arthritis; septic shock; pancreatitis; cardiac dysfunction; neuropathology; cardiac anaphylaxis; autoimmunity; genetic disease; haematopoletic disorder; platelet disorder; asthma; thrombocytopaenia; osteoporosis; severe combined immunodeficiency; allergic rhinitis; diabetes; multiple sclerosis; depression; hizheimer's disease; Parkinson's disease; neurodegenerative disorder; WO200153455-A2 neurological disorder; 22-DEC-2000; 26-JUL-2001 Homo sapiens cancer; ulcer; HIV infection; human immunodeficiency virus; 99US-0471275. 2000US-0488725. 2000US-0552317. 2000WO-US35017 SS

2001-457603/49 YT, Liu C, Drmanac

RT.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC AAM25963 The proteins can have activities based on the tissues and cells they are expressed in, such as: antinflammatory; antirheumatic; CC antiarrhritic; immunosuppressive; antipacterial; endocrine; cardiant; ccentral nervous system; virucide; anti-HIV; fungicide; antiastimatic; CC antidabetic; cytostatic; neuroprotective; antidepressant; noctropic; antiastematic; cantidabetic; cytostatic; neuroprotective; antidepressant; noctropic; CC antidabetic; cytostatic; neuroprotective; antidepressant; noctropic; CC antidabetic; noteins; and immunostimulant. The proteins and polynucleotides cencoding them can be used in gene therapy, antisense therapy and vaccine condists or antagonists of a protein and for the treatment and diagnosis of disorders associated with the activity of a protein e.g. inflammation, CC neuropathology, cardiac anaphylaxis, viral, bacterial, HIV and fungal infections, autoimmunity, genetic diseases, haematopoietic disorders, atthat, altergic canaemia, platelet disorders, thrombocytopaenia, wounds, burns, ulcers, CC altheimer's disease, Parkinson's disease, neurodegenerative and construction and rollings, severe combined immunodeficiency, eczema, allergic children's disease, Parkinson's disease, neurodegenerative and construction neuropatical disorders. Parkinson's disease, neurodegenerative and construction neuropatical disorders.
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Best Local Similarity
Matches 27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 604-605; 1217pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated human polynucleotides encoding treatment and diagnosis of e.g. cancer,
WPI; 2000-500381/45
                                                                                              26-FEB-1999;
                                                                                                                                                                                                EP1033401-A2
                                                                                                                                                                                                                                                              Human; 5' EST; expressed sequence tag; secreted protein; cDNA isolation; gene therapy; chromosome mapping; ss.
                                                                                                                                                                                                                                                                                                                                             06-OCT-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1040 BP; 194 A; 287 C; 302 G; 257 T; 0 other;
                                Dumas Milne Edwards
                                                             (GEST ) GENSET
                                                                                                                               21-FEB-2000; 2000EP-0200610
                                                                                                                                                               06-SEP-2000.
                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                Human secreted
                                                                                                                                                                                                                                                                                                                                                                                                                AAC03740 standard; cDNA; 1066 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 27; Dilarity 100.0%; Pred. No. 0.0 Conservative 0; Mismatches
                                                                                                  99US-0122487
                                                                                                                                                                                                                                                                                                              protein 5' EST, SEQ ID NO: 3738.
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                                Duclert A,
                                Giordano
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polypeptides, useful for ulcers and HIV infection
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0.04;
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The present sequence is one of a large number of 5′ ESTs derived from mRNAs encoding secreted proteins. An ORF has been identified within the sequence. The 5′ ESTs were prepared from total human RNAs or polyAr RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3′ untranslated region (UTR) of the mRNA because they are often obtained from oligo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5′ ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5′ UTR is rarely included. 5′ ESTs are derived from mRNAs with intact 5′ ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5′ ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain vectors.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID 3738; 71pp + CD-ROM; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for diagnostic, forensic, gene therapy and chromosome mapping procedures
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Search completed: November 19, Job time: 193.769 secs 2002, 03:06:38

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В δÃ

Query Match 100.0%; Score 27; I Best Local Similarity 100.0%; Pred. No.:0. Matches 27; Conservative 0; Mismatches

0; Mismatches

DB 21; .04; 0;

Length 1066; Indels

0

Gaps

0

Sequence 1066 BP; 230 A; 283 C; 310 G; 241 T; 2 other;

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Result
No.
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Maximum DB seq
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Title:
Perfect score:
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                       C
                                                                                                                            Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                          Score
   27
27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               length: 2000000000
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                                                                        Query
Match
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Listing first 45 summaries
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Copyright (c) 1993 - 2002
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em_estpl:*
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em_gss_hum:*
em_gss_inv:*
em_gss_pln:*
em_gss_vrt:*
em_gss_fun:*
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em_esthum:*
                                                                                                                                                                     em_gss_other:*
em_gss_pro:*
em_gss_rod:*
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gb_htc:*
gb_est3:*
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em_estfun:*
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 316
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377
397
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AA506459
AA503943
AA603297
AA573604
AA177005
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Compugen Ltd
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AA593245 nn07e10.s
AA506459 nh46a08.s
AA506459 nh46a08.s
AA603297 np12c11.s
AA603297 np12c11.s
AA573604 nf42d12.s
AA177005 nc03b02.s
                                                                        Description
                                                                                                                                     REFERENCE
AUTHORS
TITLE
                                                                                                                                                                                         ACCESSION
VERSION
KEYWORDS
SOURCE
ORGANISM
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AA593245
LOCUS
DEFINITION
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                                                                                                                   JOURNAL
                                                                                                                                                                                                                                                                                                                                                                    Email: cgapbs-r@mail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.,
Michael R. Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: David B. Krizman, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                  Homo sapiens

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 316)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

NATIONAL Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                       Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                        AA593245 316 bp mRNA linear EST 25-SEP-1997 nn07e10.s1 NCI_CGAP_Pr4.1 Homo sapiens cDNA clone IMAGE:1077066 similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);
                                                                                                                                                                                                                          mRNA sequence.
AA593245
AA593245.1 GI:2409007
 human.
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AA578976
2 BE840701
2 BE840537
2 BF679591
AI557591
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BF679168
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AI525089
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AI685510
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BF678386
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BF675112
BF675142
BF679394
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BF678076
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BF678076
BF67376
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BF6737323
BF673233
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                                                                                                                                                                                                                                                                                                                                     ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                             AA533652 nj72908.s
AI68510 tu36b10.x
BI046475 MR3 FW020
AI525128 promrna-7
AA603329 np15b08.s
AA573575 nf41f10.s
BE840533 RC1-FN018
BE840641 RC1-FN018
AA639901 np08f03.s
BE840701 RC1-FN018
BE840777 RC1-FN018
BE840777 RC1-FN018
BE840777 RC1-FN018
BE777591 60215403
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BF677591 pt2.1-1.F
BF677951 602267731
AI547785 PN001_AH_
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BF677366 602135902
BF677376 602085123
BF677376 602085123
BF677376 602085123
BF677376 602085123
BF677376 602085123
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Euteleostomi;
Homo.

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FEATURES
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VERSION
KEYWORDS
SOURCE
ORGANISM
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AA506459
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ORIGIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AUTHORS
TITLE
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Best Local
                                                                                                                                                                 source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tumor Gene Index
Unpublished (1997)
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Flissue Procurement: David G. Bostwick, M.D., Rodrigo F. Chuaqui,
M.D., Michael R. Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: David B. Krizman, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bb.llnl.gov/Dbrp/image/Image.html
Insert Length: 442 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    l Similarity 100 27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AA506459 a36 bp mrNA linear EST 20-AUG-1997 nh46a08.sl NCI_CGAP_Pr5 Homo sapiens cDNA clone IMAGE:955382 similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 336)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AA506459
AA506459.1 GI:2242699
EST.
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High quality sequence stop: 293.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mRNA sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /lab_host="DH10B"
/lab_host="DH10B"
/note="Organ: prostate; Vector: pamp10; mrNA made from /note="Organ: prostate; intraepithelial neoplasia (high-grade), cDNA made by oligo-dT priming. Non-directionally cloned.
Size-selected on agarose gel, average insert size 600 bp.
CDNA Library Preparation: David B. Krizman, ph.D.
Reference: Krizman et al. (1996) Cancer Research, ph.D.
56:5380-5383. cDNA Library Arrayed by: Greg Lennon, ph.D.
DNA Sequencing by: Washington University Genome Sequencing
Center
              /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:955382"
/clone_lib="NCI_CGAP_Pr5"
/sex="male"
/tissue_type="prostate"
/lab_host="DH10B"
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grade"
                                                                                                                                                                               Location/Qualifiers
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/clone_lib="NCI_CGAP_Pr4.1"
/sex="male"
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/db_xref="taxon:9606"
/note="Vector: pampl0; mRNA made from normal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
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ORIGIN
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AUTHORS
TITLE
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ORGANISM
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ORIGIN
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                                                                                182
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Query Match
Best Local S
Matches 27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                189 GTTCTGGTGCACCCCAGTGGGTCCTC 215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 GTTCTGGTGCACCCCCAGTGGGTCCTC 27
                                               1 GTTCTGGTGCACCCCCAGTGGGTCCTC 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tumor Gene Index

Oppublished (1997)

Email: cgapbs-remail.nih.gov

Email: cgapbs-remail.nih.gov

Tissue Procurement: David G. Bostwick, M.D., Rodrigo F. Chuaqui,

M.D., Michael R. Emmert-Buck, M.D., Ph.D.

cDNA Library Preparation: David B. Krizman, Ph.D.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Mashington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 454 Std Error: 0.00

Seq primer: -40ml3 fwd. ET from Amersham

High quality sequence stop: 365.

Std Error: 0.00

Seq primer: -40ml3 fwd. ET from Amersham

Location/Qualifiers
l Similarity
27; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA503943

AA503943

371 bp mRNA linear EST 20-AUG-1997

nh38d05.sl NCI_CGAP_Pr5 Homo sapiens cDNA clone IMAGE:954633

similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EST
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ilarity 100.0%;
Conservative (
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                                                                                                                                                                                                             /tissue_type="prostate"
/lab_host="DH10B"
/lab_host="bH10B"
/lab_host="bH10B"
/note="Vector: pampl0; mRNA made from normal prostatic note="Vector: pampl0; mRNA made by oligo-dT priming.

kon-directionally cloned. Size-selected on agarose gel, average insert size 600 bp. "

a 112 c 111 g 85 t
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Non-directionally cloned. Size-selected on agarose gel,
average insert size 600 bp. "
                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:954633"
/clone_lib="NCI_CGAP_Pr5"
/sex="male"
                                                                                              0;
                                                                                              Score 27; DB 9;
Pred. No. 0.9;
; Mismatches 0
                                                                                                 0,
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                                                                                                                                          Length 371;
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RESULT 5
AA573604
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Best Local Similarity
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                     ORGANISM
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                                                                                                                                                                                                                                                                                           288
                                                                                                                                                                                                                                                                                                                               GTTCTGGTGCACCCCCAGTGGGTCCTC 27
                                                                                                                                                                                                                                                                                         GTTCTGGTGCACCCCCAGTGGGTCCTC 262
                                                                                                                                                                                                                                                                                                                                                                        27;
                                                            EST.
                                                                                                                                         AA573604 397 bp mRNA linear EST 12-SEP-1997 nf42d12.sl NCI_CGAP_Pr2 Homo sapiens cDNA clone IMAGE:916439 similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      www-bio.llnl.gov/bbrp/image/image.html.
Insert Length: 449 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 345.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AA603297 377 bp mRNA linear EST 21-OCT-1997 np12c11.s1 NCI_CGAP_Pr3 Homo sapiens cDNA clone IMAGE:1116116 similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);,
                   Homo sapiens
                                                                                 AA573604.1 GI:2348132
                                                                                                    mRNA sequence.
AA573604
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cDNA Library Preparation: David B. Krizman, Ph.D. cDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D., Michael Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                          human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                    100.0%; ilarity 100.0%; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note-"Vector: pAMP10; Site_1: Not1; Site_2: EcoRI; 1st strand cDNA was primed with oligo(dT)17 on 50 ng of DNAse-treated, total cellular RNA obtained from 5,000-10,000 microdissected cells histologically-determined to be fully malignant prostate cancer cells. Double-stranded cDNA was ligated to EcoRI adaptors, 5 cycles of PCR applied to the cDNA with an adaptor-specific primer, and the resulting PCR product subcloned into pAMP10 by the UDG-cloning method (Life Technologies). Average insert size is 600 bp. NOTE: Not directionally cloned. This library was constructed by David Krizman."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:9606"
/clone="IMAGE:1116116"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone_lib="NCI_CGAP_Pr3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /dev_stage="45 years old"
/lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                                                                                    Score 27; DB 9
Pred. No. 0.9;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                               Length 377;
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COMMENT
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AUTHORS
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AA177005
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Best Local Similarity
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                                                                                 Tumor Gene Index
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                                                                                                                                                                                                                           numan
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Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.
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95 GTTCTGGTGCACCCCCAGTGGGTCCTC 121
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Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D.,
Michael Emmert-Buck, M.D., Ph.D.
                                                         Unpublished (1997)
                                                                                                             NCI-CGAP http://www.ncbi.nlm.nih.góv/ncicgap
National Cancer Institute, Cancer Genome Ana
                                                                                                                                                                       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 435)
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High quality sequence stop: 350.
Location/Qualifiers
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Unpublished (1997)
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ligated to EcoRI adaptors, 5 cycles of PCR applied to the cDNA with an adaptor-specific primer, and the resulting PCR product subcloned into pawP10 by the UDG-cloning method (Life Technologies). Average insert size is 600 bp. NOTE: Not directionally cloned. This library was constructed by David Krizman."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note="Vector: pAMP10; Site_1: Not1; Site_2: EcoRI; 1st strand cDNA was primed with oligo(dT)17 on 50 ng of DNAse-treated, total cellular RNA obtained from 5,000-10,000 microdissected preneoplastic cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               histologically-determined to be prostatic intraepithelial neoplasia 2 (PIN2) cells. Double-stranded cDNA was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone="IMAGE:916439"
/clone_lib="NCI_CGAP_Pr2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /dev_stage="45 years old"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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0.9;
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                                                                                                                Anatomy Project (CGAP),
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AA533652
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VERSION
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Best Local
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                                                                                                                                     Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo F. Chuaqui,
M.D., Michael R. Emmert-Buck, M.D., Ph.D.
CDNA Library Preparation: David B. Krizman, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
cDNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortimm/LLNL at:
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cDNA Library Preparation: David B. Krizman, Ph.D.

cDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D.

cDNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be

found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

Seq primer: -40M13 fwd. from Amersham.

Location/Qualifiers
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A53 bp mRNA linear EST 21-AUG-1997
n)72g08.sl NCI_CGAP_Pr10 Homo sapiens cDNA clone IMAGE:998078
similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27; Conserv
Trace considered overall poor quality Insert Length: 1654 Std Error: 0.00 Seg primer: -40ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                 Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 453) NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap. Nclional Cancer Institute, Cancer Genome Anatomy Project (CGAP), Mational Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mRNA sequence
AA533652
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                                                                                                                 /ww-bio.llnl.gov/bbrp/image/image.html
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54 a 119 c 146 g 105 t 1 others
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/db_xref="taxon:9606"
/clone="IMAGE:302"
/clone_lib="NCI_CGAP_Pr2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /dev_stage="45 years old"
/lab_host="DH10B"
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Pred. No. 0.9;
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AI685510/c
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Emmail: cgapbs-r@mail.nih.gov

Tissue Procurement: Michael J. Brownstein, M.D., Ph.D., Michael R.

Emmert-Buck, M.D., Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D.

CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be

found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

Seq primer: -40UP from Gibco.
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AI685510
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Location/Qualifiers
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Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 454)

NCI-GGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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/dev_stage="adult"
//lab_host="hylloB"
//lab_h
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:2253115"
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                                                                                                                                                                                                                                                                                                                                                                                                          /clone_lib="NCI_CGAP_Pr28"
/sex="male"
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/db_xref="taxon:9606"
/clone="IMAGE:998078"
/clone_lib="NCI_CGAP_Pr10"
/sex="male"
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/lab_host="DH10B"
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323
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               1 GTTCTGGTGCACCCCCAGTGGGTCCTC 27
GTTCTGGTGCACCCCCAGTGGGTCCTC 349
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MR3-FN0209-070201-010-b11 FN0209
BI046475
                                                                                                                                                                                                                                                                                                                                                                                                                  Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?t1=MR3&t2=MR3-FN0209-
070201-010-b11&t3=2001-02-07&t4=1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Dias Neto,E., García Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Laboratory of Cancer Genetics Ludwig Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Shotgun sequencing of the human transcriptome with ORF expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Simpson, A.J.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Contact: Simpson A.J.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence tags
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (bases
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                                                                                                                                            83
                                                              Conservative
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                                                                                                                                         /note="Organ: prostate_normal; Vector: puc18; Site_1: SmaI; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."

a 134 c 144 g 103 t 1 others
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                                                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone_lib="FN0209"
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                                                                            100.0%;
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                                                                                                                                                                                                                                         VERSION
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui,
, Michael Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: David B. Krizman, Ph.D.
cDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph
                                                                                                                                                                                                                                                                                                                                                               AA603529 484 bp mRNA linear EST 08-OC np15b08.s1 NCI_CGAP_Pr3 HOMO sapiens cDNA clone IMAGE:1116375 similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUM
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Department of Molecular Biotechnology,
Washington, Seattle, WA 98195
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AI525128
AI525128.1 GI:4439263
                                                                                                   Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                     NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy
                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 484)
                                                                                                                                                                                                                                                Homo sapiens
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AA603529
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Huang,G.M., Ng,W.l., Farkas,J.,
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Mammalia; Eutheria;
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llarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Washington).
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/clone_lib="bvtumor"
/note="Organ: Prostate; Vector: pBluescript; DcDNA library was constructed using Lambda ZP I (Stratagene). mRNA was extracted from a fresh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tumor tissues (Urology Department, University of
Washington) "
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/db_xref="taxon:9606"
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                                                                                                                                                         Project (CGAP),
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Ph.D.

source

/clone\_lib= /sex="Male"

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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Edikaryota; Metazoa; Chordata; Catarrhini; Hominidae; Homo.

Ediages 1 to 485)

RNI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

RNI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

RNI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

RNI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

Timor Gene Index

Unpublished (1997)

Contact: Robert Strausberg, Ph.D.

Email: cgapbs-remail.nih.gov

Fissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D., Michael Emmert-Buck, M.D.; Ph.D.

CDNA Library Preparation: David B. Krizman, Ph.D.

CDNA Library Preparation: David B. Krizman, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center found through the I.M.A.G.E. Consortium/LINL at:

www-bio.llnl.gov/bbrp/image.lmage.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Trace considered overall poor quality Insert Length: 1535 Std Error: 0.00 Seq primer: -40ml3 Fwd. Er from Amersham High quality sequence stop: 1.

Location/Qualifiers
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AA573575
AA5713575
Inf41F10.s1 NCI_CGAP_Pr2 Homo. Sapiens cDNA clone IMAGE:916363
similar to 9b:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);,
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1 Similarity 100.0%;
27; Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               //note-/vector: pAMPIO; Site_1: Not1; Site_2: EcoRI; 1st strand cDNA was primed with oligo(dT)17 on 50 ng of .00-10 NASe-treated, total cellular RNA obtained from 5,000-10 .000 microdissected cells histologically-determined to be fully malignant prostate cancer cells. Double-stranded cDNA was ligated to EcoRI adaptors; 5 cycles of PCR applied to the cDNA with an adaptor-specific primer, and UDG-cloning method (Life Technologies). Average insert size is 600 bp. NOTE: Not directionally cloned. This library was constructed by David Krizman."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:1116375"
/clone_lib="NCI_CGAP_Pr3"
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/lab_host="DH10B"
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Pred. No. 0.91;
Mismatches (
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COMMENT
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AUTHORS
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SOURCE
ORGANISM
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BE840533/c
LOCUS
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Best Local :
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Tel: +55-11-2707001

Tex: +55-11-2707001

Email: asimpson@ludwig.org.br

This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL (http://www.ludwig.org.br/scripts/gethtml2.pl?ti=st2=RC1-FN0188-260 700-022-f066t3=2000-07-266t4=1)

Seq primer: puc l8 forward High quality sequence stop: 495.

Location/Qualiflers

1. 495
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Eukaryota, Metazoa, Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota, Metazoa; Chordata; Catarrhini; Hominidae; Homo.

| (bases 1 to 495)
Dias Neto,E., Garcia Correa,R., Verjovski-Almeidae,S., Eriones,M.R.,
Nagal,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Soldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,
Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tch 100.0%; sal Similarity 100.0%; Fal Similarity 00.0%; Fallow 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence tags
Proc. Natl. Acad. Sci. U.S.A. 97 (7),
20202663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Simpson, A.J.
Shotgun sequencing of the human transcriptome with ORF expressed
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/organism="Homo sapiens"
/organism="Homo sapiens"
/organism="Image:91363"
/clone_lib="NCI_CGAP_Pr2"
/sex="Male"
NCI_CGAP_Pr2"
/sex="Male"
NCI_CGAP_Pr2"
/sex="Male"
/olone_lib="NCI_CGAP_Pr2"
/sex="Male"
NCI_CGAP_Pr2"
/sex="Male"
/
/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone_lib="FN0188"
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Pred. No. 0.91;
Mismatches 0
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0

FEATURES

REFERENCE AUTHORS TITLE

ACCESSION VERSION KEYWORDS SOURCE ORGANISM

Homo sapiens

mRNA sequence. AA573575 AA573575.1 GI:2348103

RESULT 12 AA573575 LOCUS DEFINITION

187

ORIGIN BASE COUNT

68

Query Match Best Local S Matches 27

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REFERENCE
AUTHORS
BASE COUNT
ORIGIN
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BE840441/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?t1=&t2=RC1-FN0188-210
700-021-a02&t3=2000-07-21&t4=1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 505)
Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shotgun sequencing of the human transcriptome with ORF expressed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Natl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000)
               application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."

164 c 136 g 78 t
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a 161 c 131 g 77 t 1 others
                                                                                                                                           /note="Organ: prostate_normal; Vector: puc18; Site_1: SmaI
; Site_2: SmaI; A mini-library was made by cloning
products derived from ORESTES PCR (U.S. Letters Patent
                                                                                                                                                                                                                                                  /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone_lib="FN0188"
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                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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Query Match

Score 27;

DВ 12;

Length 505;

Search completed: November Job time: 1545.46 secs

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SOURCE
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                                                                                Matches
                                                                                                                          Query Match
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  187
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                                                                                                    Local
                 1 GTTCTGGTGCACCCCCAGTGGGTCCTC
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                                                                                27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              www-bio.llnl.gov/bbrp/image/image.html
Seg primer: -40m13 fwd. ET from Amersham
High quality sequence stop: 367.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D., Michael Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Tumor Gene Index Unpublished (1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mammalia; Eutheria;
1 (bases 1 to 523)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
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                                                                                  Conservative
                                                                                                                                                                            /note="Vector: pAMP10; Site_1: Not1; Site_2: EcoRI; 1st strand cDNA was primed with oligo(dT)17 on 50 ng of DNAse-treated, total cellular RNA obtained from 5,000-10,000 microdissected cells histologically-determined to be fully malignant prostate cancer cells. Double-Stranded cDNA was ligated to EcoRI adaptors, 5 cycles of PCR applied to the cDNA with an adaptor-specific primer, and the resulting PCR product subcloned into pAMP10 by the UDG-cloning method (Life Technologies). Average insert size is 600 bp. NOTE: Not directionally cloned. This library was constructed by David Krizman."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/db_xref="taxon:9606"
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RESULT 2 US-09-100-264-4

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APPLICANT: Young, Charles Y.F.
APPLICANT: Tindall, Donald J.
APPLICANT: Klee, George G.
TITLE OF INVENTION: Method for detection of breast cancer
FILE REFERENCE: 545.0060IS1
CURRENT APPLICATION UNUMEER: US/09/100.264
CURRENT FILING DATE: 1998-06-19
CURRENT APPLICATION NUMBER: US 60/050.963
EARLIER APPLICATION NUMBER: US 60/050.963
EARLIER FILING DATE: 1997-06-20
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LENGTH: 711
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Patent No. 6235486
GENERAL INFORMATION:
                                                                                                                                                                                             COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/358,782D
FILING DATE: 15-DEC-1994

CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Beardell, Lori Y.
REGISTRATION NUMBER: 34,293
REFERENCE/DOCKET NUMBER: 31.297

TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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TELECOMMUNICATION INFORMATION:
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                                                                                              TELEPHONE: 215-568-3100
TELEFAX: 215-568-3439
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 992 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Mulholland, S. Grant
APPLICANT: Moreno, Jose
APPLICANT: Fischer, Rainer
TITLE OF INVENTION: Methods of Detecting Micrometastasis of Prostate
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Moodcock Washburn Kurtz Mackiewicz &
STREET: One Liberty Place 46th. Floor
CITY: Philadelphia
STRIE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Gomella, Leonard
APPLICANT: Mulholland, S. Grant
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                         TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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DNA (genomic)
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  В
                                                                                                                                                MOLECULE TYPE: DN
HYPOTHETICAL: NO
ANTI-SENSE: NO
US-08-764-527A-13
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US-08-764-527A-13
                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/358,782

FILING DATE: 15-DEC-1994

ATTORNEY/AGENT INFORMATION:

RAME: Beardell, Lori Y.

REGISTRATION NUMBER: 34,293

REGISTRATION NUMBER: TUU-1327

TELEPANONE: 215-568-3100

TELEPAN: 215-568-310

TELEPAN: 215-568-310

TELEPAN: 215-568-310

TELEPAN: 215-568-310

TELEPAN: 215-568-310

INFORMATION FOR SEQ ID NO: 13:

SEQUENCE CHARACTERISTICS:

LENGTH: 992 base pairs

TYPE: DIGITAL CALLERY SEGUENCE: 1515-611
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Best Local Similarity 100.
Matches 27; Conservative
                                                                       Query Match
Best Local Similarity
Matches 27; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURENUT APPLICATION DATA:
APPLICATION NUMBER: US/08/764,527A
FILING DATE:
CLASSIFICATION:
CLASSIFICATION:
CLASSIFICATION:
CLASSIFICATION:
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CORRESPONDENCE ADDRESS: Hoodcock Washburn Kurtz Mackiewicz & STREET: One Liberty Place 46th. Floor CITY: Philadelphia
STATE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Fischer, Rainer
TITLE OF INVENTION: Methods of Detecting Micrometastasis
TITLE OF INVENTION: Prostate
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1 GTTCTGGTGCACCCCCAGTGGGTCCTC 27
                                                                                                                                                                                                                                          STRANDEDNESS: single
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                                                                         100.0%; Score 27; I ilarity 100.0%; Pred. No. 0 Conservative 0; Mismatches
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RESULT 5 US-09-605-785-616; A)

Application US/09605785

APPLICANT: Xu, Jiangchun

Dillon, Davin C.

Mitcham, Jennifer L.

INFORMATION:

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; SOFTWARE: FastSEQ for Windows Version 3.0; SEQ ID NO 616; LENGTH: 1350; TYPE: DNA; ORGANISM: Homo sapien US-09-605-785-616
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US-08-358-782D-14
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Patent No. 5
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CURRENT FILING DATE: 2000-06-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Hepher, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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            ATTORNEY/AGENT INFORMATION:
NAME: BEARDELL LOTI Y.
REGISTRATION NUMBER: 34,293
REFERENCE/DOCKET NUMBER: TJU
                                                                             SOFTWARE: PatentIn Release #1.0, VGCURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/358,782D
FILING DATE: 15-DEC-1994
CLASSIFICATION: 435
                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                          APPLICANT: Fischer, Rainer
TITLE OF INVENTION: Methods of Detecting Micrometastasis of Prostate
NUMBER OF SEQUENCES: 14
TELECOMMUNICATION INFORMATION:
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STATE: F.,
STATE: 19103
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                                                                                                                                                                                                                                                                       CITY: Philadelphia
                                                                                                                                                                                                                                                                                         ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & STREET: One Liberty Place 46th, Floor
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27; Conserv
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Fanger, Gary R.
Retter, Marc W.
Stolk, John A.
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100.0%; Pred. No.
tive 0; Mismatch
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              TJU-1327
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                                                                                                                                                   Version #1.25
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0.0072;
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; MOLECULE TYPE: I
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
US-08-358-782D-14
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US-08-764-527A-14
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                                                                                                            TELEFAX: 215-568-3439
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO:
                                                                                                                                                         REFERENCE/DOCKET NUMBER: TO TELECOMMUNICATION INFORMATION TELEPHONE: 215-568-3100
                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08
FILING DATE: 15-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: Beardell, Lori Y.
                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
            HYPOTHETICAL:
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                        MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
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STATE: r..
19103
                                                                  LENGTH: 1462 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                   REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
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                                                         TOPOLOGY:
                                                                                              LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Philadelphia
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                                                       linear
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                                          DNA (genomic)
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100.0%; Pred. No. 0.
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Query Match Best Local Similarity

100.0%;

Score Pred.

27; No.

DB 2; 0.0073;

Length 1462;

В QΥ

181

COUNTRY:

Matches

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; NAME/KEY:
; LOCATION:
US-08-844-024-1
                                                                                      RESULT 9
US-08-718-547-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:

APPLICATION UNMER: 08/229,391
APPLICATION UNMER: 08/229,391
FILING DATE: 15-APR-1994
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION UNMBER: 28,678
REFERENCE/DOCKET NUMBER: 43677/J
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 977-9550
TELEEX: 42823 COOP UI
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 1729 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Application US/08844024 Patent No. 5840494
                                Sequence 1, Application US/08718547 Patent No. 5976794 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Katz, Aaron E., et al.
TITLE OF INVENTION: A Method For Molecular Staging Of
TITLE OF INVENTION: Prostate Cancer
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPALIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE:
ORIGINAL SOURCE:
APPLICANT: Katz, Aaron E.; Buttyan, Ralph; Raffo, Anthony; APPLICANT: Olsson, Carl A.
                                                                                                                                                         STRANDEDNESS: SI
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/844,024 FILING DATE: 18-APR-1997
                                                                                                                                                                                                                                                                                                                                                                        ORGANISM:
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ADDRESS:

Cooper 6 Dunham

COOPER 7 DUNHAM

TATE: New York

TATE: New York
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27; Conserv
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378..1088
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Pred. No. 0.0
0; Mismatches
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                                  NUMBER OF SEQ ID NOS: 35
SCPTWARE: FastSEQ for Windows Version 3.0; SEQ ID NO 31
I LENGTH: 7130
TYPE: DNA
CORGANISM: Homo sapiens
US-09-056-105-31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY:
; LOCATION:
US-08-718-547-1
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US-09-056-105-31
                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                      Sequence 31, Application US/09056105 Patent No. 6287569
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                         APPLICANT: KIPPS, THOMAS J.

APPLICANT: WU, YUNGI WITH ENHANCED INTRACELLULAR
TITLE OF INVENTION: VACCINES WITH ENHANCED INTRACELLULAR
TITLE OF INVENTION: PROCESSING
FILE OF INVENTION SPROCESSING
FILE REFERENCE: 33/221
CURRENT APPLICATION NUMBER: US/09/056,105
CURRENT FILING DATE: 1998-04-06
EARLIER APPLICATION NUMBER: 60/04/3,467
EARLIER FILING DATE: 1997-04-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX:
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENCTH: 1729 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
CLASSIFICATION: 435
ATTORNEY, AGENT INFORMATION:
NAME: White, John P.
REGISTRATION UNMBER: 28,678
REFERENCE/DOCKET NUMBER: 4367
TELECOMMUNICATION INFORMATION:
TELEPIONE: (212) 278-0400
TELEPIAX: (212) 391-0525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LEET: 1. COOPER & Dunham LLP
LIRES Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 1036
TOMPUTPPD 1036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
COMPUTER: IEM PC compatible
COPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
COPERATE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/718,547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS
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IGINAL SOURCE:
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27; Conserv
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PE: cDNA
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378..1088
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    100.0%;
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      Score 27;
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Pred. No. 0.
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0.0074;
es 0;
      4;
      Length 7130;
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Matches Query Match

Best Local Similarity Matches 27; Conserv

Conservative

100.0%; Pred. No. 0.0082; Nigmatches 0;

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US-08-622-046B-13
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US-08-622-046B-2
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                                                                                                                                           Sequence 13, Application US/08622046B Patent No. 6103237
                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                             GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Stable Variant NUMBER OF SEQUENCES: 21
                                                                                               APPLICANT: Saedi, Mohamn APPLICANT: Mikolajczyk,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: 612-373-6903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Saedi, Mohamma APPLICANT: Mikolajczyk,
                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Stable variant HK2 Polypeptide NUMBER OF SEQUENCES: 21
                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: 11 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Woessner, Warren D.
REGISTRATION NUMBER: 30,440
REFERENCE/DOCKET NUMBER: 476.001US1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: FILING DATE: 26-MAR CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                   85 GTCCTGGTGCACCCCCAGTGGGTGCTC 111
                                                                                                                                                                                                                                                                                                       Local
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               CITY:
                            STREET:
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                                       ADDRESSEE:
                                                                                                                                                                                                                                                1 GTTCTGGTGCACCCCCAGTGGGTCCTC 27
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25; Conservative
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                            P.O.
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                                                                                                                Saedi, Mohammed S
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                                                                                                                                                                                                                                                                                                                                                                               linear
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                            Schwegman,
O. Box 2938
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                                                                                                                                                                                                                                                                                                       88.1%;
92.6%;
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Pred. No. 0.15;
0; Mismatches
                                          Woessner & Kluth, P.A.
                                                                                   HK2 Polypeptide
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                                                                                                                                                                                                                                                                                                                    DB 3;
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Query Match
Best Local Similarity
Watches 25; Conserve
RESULT 14
US-08-768-859A-7
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                                                                                                                                                                                             ; TYPE: DNA; ORGANISM: Homo sapiens US-09-100-264-2
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US-09-100-264-2
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                                                                                                                                                                                                                                           SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 2
LENGTH: 711
                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence 2, Application US/09100264
Patent No. 6235486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Young, Charles Y.F.
APPLICANT: Tindall, Donald J.
APPLICANT: Klee, George G.
TITLE OF INVENTION: Wethod for detection of
FILE REFERENCE: 545.006US1
                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/100,264
CURRENT FILING DATE: 1998-06-19
EARLIER APPLICATION NUMBER: US 60/050,963
EARLIER FILING DATE: 1997-06-20
                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 612-339-3061
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: 15 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 711 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: 612-373-6903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 30,440
REFERENCE/DOCKET NUMBER: 476.001US1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FILING DATE: 26-MAI
                                                                85 GTCCTGGTGCACCCCCAGTGGGTGCTC 111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 GTTCTGGTGCACCCCCAGTGGGTCCTC 27
                                                                                 1 GTTCTGGTGCACCCCCAGTGGGTCCTC 27
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25; Conserv
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DEDNESS: double
                                                                                                                              Conservative
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92.6%;
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92.6%;
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Pred. No. 0.
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Sequence 7, Application US/08768859A Patent No. 6013471
GENERAL INFORMATION:

APPLICANT:

Tindall, Donald J. Young, Charles Y.F.

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STATE.

COUNTRY: USA
ZIP: 55402

ZIP: 55402

COMPUTER READABLE FORM:

MEDJIM TYPE: Floppy disk

COMPUTER: IBM PC COMPATIBLE

SOFTWARE: PC-DOS/MS-DOS

SOFTWARE: PC-DOS/MS-DOS

SOFTWARE: PC-DOS/MS-DOS

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/768,859A

FILING DATE: 17-DECEMBER-1995

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

REFERENCE/DOCKET NUMBER: 30,440

REFERENCE/DOCKET NUMBER: 355.002US3

TELECOMMUNICATION INFORMATION:

TELEPHONE: 612-339-3061

TELEPHONE: 612-339-3061

TELEPHONE: 612-339-3061

TELEPHONE: 612-339-3061

TELEPHONE: 612-339-3061
          Sequence 7, Application US/08767820A

Fatent NO. 6093796

GENERAL INFORMATION.

APPLICANT: Tindall, Donald J.

APPLICANT: Tindall, Donald J.

APPLICANT: Saedi, Mohammed S.

ITTLE OF INVENTION: DNA Encoding HK2 Variant Polypeptides

NUMBER OF SEQUENCES: 21

CORRESPONDENCE ADDRESS:

ADDRESSE: Schwegman, Lundberg, Woessner & Kluth, P.A.

STREET: PO. Box 2938

CITY: Minneapolis

STATE: MN

COUNTRY: USA
ITPLE: 5402

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION UNBER: US/08/767,820A

FILING DATE: 17-DECEMBER-1996

CLASSIFICATION. 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; NAME/KEY: CDS; LOCATION: 7..720
US-08-768-859A-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 15
US-08-767-820A-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPAX: 612-339-3001
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 760 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 92.6
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Saedi, Mohammed S.
TITLE OF INVENTION: DNA Encoding HK2 Variant Polypeptides
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: P.O. Box CITY: Minneapolis
STATE: MN
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Schwegman, Lundberg, Woessner & Kluth, P.A. P.O. Box 2938
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88.1%; Score 23.8; DB 3; Length 760; 92.6%; Pred. No. 0.15; tive 0; Mismatches 2; Indels
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; NAME/KEY: CDS
; LOCATION: 7..720
US-08-767-820A-7
Search completed: November 19, 2002, 05:20:21 Job time: 53.9615 secs
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                                                                                                  Qy
                                                                                                                          Query Match 92.6
Best Local Similarity 92.6
Matches 25; Conservative
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TELEFAX: 612-339-3061
INFORMATION FOR SEQ ID NO: 7:
                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
AMME: WOESSNEY, WARTEN D.
REGISTRATION NUMBER: 30,440
REFERENCE/DOCKET NUMBER: 545.002US2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 612-339-0331
                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 760 base pairs
                                                                                                                                                                                                                                                                   FEATURE:
                                                                                                                                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: cDNA
                                                                       94 GTCCTGGTGCACCCCAGTGGGTGCTC 120
                                                                                                                                                                                                                                                                                                                  LENGTH: 760 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                          1 GTTCTGGTGCACCCCCAGTGGGTCCTC 27
                                                                                                                                                         92.6%;
                                                                                                                                         Score 23.8; DB 3;
Pred. No. 0.15;
0; Mismatches 2;
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                                                                                                                                             Indels
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Gaps

0,

GenCore version 5.1.3 Copyright (c) 1993 - 2002 Compugen Ltd

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Result
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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DB
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1: //gqn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq:*

2: //cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*

3: //gqn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*

4: //gqn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*

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12: //gqn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*

13: //gqn2_6/ptodata/1/pubpna/US00_NEW_PUB.seq:*

14: //gqn2_6/ptodata/1/pubpna/US00_NEW_PUB.seq:*

14: //gqn2_6/ptodata/1/pubpna/US00_NEW_PUB.seq:*
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         US-09-829-004A-8
US-09-789-143-616
US-09-780-669-616
US-09-822-827-616
US-09-862-827-828-7-98
US-09-822-827-980
US-09-755-100-5
US-09-755-100-1
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US-09-755-300-153
US-09-925-300-153
US-09-925-300-814
US-09-925-300-814
US-09-822-827-790
US-09-822-827-790
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111.898 Million cell updates/sec
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Sequence 8, Appli
Sequence 616, App
Sequence 616, App
Sequence 518, App
Sequence 977, App
Sequence 980, App
Sequence 5, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 2, Appli
Sequence 1, Appli
Sequence 2307, App
Sequence 2161, App
Sequence 2101, App
Sequence 2307, App
Sequence 790, App
Sequence 790, App
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Sequence 790, App
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1248	1248	1215	1215	1215	1214	1214	1214	1214	1167	1167	1167	1167	1167	1119	1119	1119	1119	1119	871	765	765	765	234	234	234
10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10
	US-09-759-143-171	822-827	-780-669	759-143		-822-827	-09-780-669	-09-759-143	-115-453	-822-827	-030	-09-780-669	-09-759-143	US-09-115-453-177	-822-827-	-09-030-606	-780-669-	US-09-759-143-177	US-09-956-999-7	-822-827	-780-669	-759-143	-827	-780-669-3	US-09-759-143-328
	Sequence 171, App	326,	326,	326,	225,	225,	225,	225,	175,	175,	175,	175,	175,	177,	177,	177,	177,	17:	7,	52	52,	52	328		Sequence 328, App

## ALIGNMENTS

RESULT 1 US-09-829-004A-8

Sequence 8, Application US/09829004A Patent No. US20020132976A1 GENERAL INFORMATION:

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RESULT 2
US-09-759-143-616
                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: CLP314 US-09-829-004A-8
              Sequence 616, Application US/09759143
PATENT NO. US20020022248A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/829,004A
CURRENT FILING DATE: 2001-04-16
PRIOR APPLICATION NUMBER: US 60/195,456
PRIOR FILING DATE: 2000-04-10
NUMBER OF SEQ ID NOS: 12
SOFTWARE: Patentin version 3.0
SEQ ID NO 8
LENGTH: 27
                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Chong, Pele
APPLICANT: Pedyczak, Artur
APPLICANT: Sia, Charles Dwo Yuan
TITLE OF INVENTION: Immunogenic Peptides Derived from Prostate-Specific Antigen
TITLE OF INVENTION: Uses Thereof
FILE REFERENCE: 11014-24
   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
                                                                                                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                     Xu, Jiangchun
Dillon, Davin (
Mitcham, Jenni:
                                                                                                                                                                                                                                     100.0%; ilarity 100.0%; Conservative 0
Davin C.
Jennifer L.
                                                                                                                                                                                                                                        0;
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                                                                                                                                                                                                                                        Mismatches
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Harlocker,

Susan L. Robert A.

Jiang, Yuqui Henderson,

```
APPLICANT: Vedvick, Thomas S.
APPLICANT: Carter, Darrick
APPLICANT: Li, Samuel
APPLICANT: Wang, Aijun
APPLICANT: Wang, Aijun
APPLICANT: Welky, Yasir A.W.
APPLICANT: Hepler, William
APPLICANT: Houghton, Raymond L.
APPLICANT: Houghton, Raymond L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 21012.437C24
CURRENT APPLICATION NUMBER: US/09/780,669
CURRENT APPLICANTED ANTE: 2001-02-09
NUMBER OF SEQ ID NOS: 943
SOPTWARR: FastSEQ for Windows Version.3.0
SEQ ID NO 616
SEQ ID NO 616
LENGTH: 1350
TYPE: DNA
ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Retter, Marc W.
APPLICANT: Stolk, John A.
APPLICANT: Day, Craig H.
APPLICANT: Vedvick, Thomas S.
APPLICANT: Vedvick, Thomas S.
APPLICANT: Vedvick, Thomas S.
APPLICANT: Li, Samuel
APPLICANT: Wang, Aijun
APPLICANT: Wang, Aijun
APPLICANT: Wang, Aijun
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C23
CURRENT APPLICATION NUMBER: US/09/759,143
CURRENT FILING DATE: 2001-01-12
NUMBER OF SED ID NOS: 934
SOFTWARE: FastSED for Windows Version 3.0
SED ID NO 616
LENGTH: 1350
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 3

US-09-780-669-616

; Sequence 616, Application US/09780669

parent No. US20020051977A1
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APPLICANT:
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-09-759-143-616
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27; Conserv
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Vedvick, Thomas S.
Carter, Darrick
Li, Samuel
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Fanger, Gary R.
Retter, Marc W.
Stolk, John A.
Day, Craig H.
Vedvick, Thomas S.
Carter, Darrick
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Fanger, Gary R.
Retter, Marc W.
Stolk, John A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Harlocker, Susan L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Henderson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mitcham, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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RESULT 6

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RESULT 4

US-09-822-827-616

US-09-822-827

Sequence 616, Application US/09822827

Patent No. US2002081680A1

GENERAL INFORMATION:

APPLICANT: XU, Jiangchun

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER

FILE REFERENCE: 210121.534C1

CURRENT APPLICATION NUMBER: US/09/822,827

CURRENT FILING DATE: 2001-03-28

NUMBER OF SEQ ID NOS: 982

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 616

LENGTH: 1350
                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/969,708
CURRENT FILING DATE: 2001-10-03
PRIOR APPLICATION NUMBER: US/60/237,606
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: US/60/237,608
PRIOR APPLICATION NUMBER: US/60/237,425
PRIOR FILING DATE: 2000-10-03
NUMBER OF SEQ ID NOS: 658
SOPTWARE: Patentin version 3.0
SEQ ID NO 598
LENGTH: 1729
TYPE: DNA
ORGANISM: Homosapiens
US-09-969-708-598
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 598, Application US/09969708
Patent No. US20020102532A1
GENERAL INFORMATION:
APPLICANT: Augustus, Meena
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Sign
TITLE OF INVENTION: Sets
FILE REFERENCE: 689290-70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
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US-09-969-708-598
                                                                         Query Match
Best Local S
Matches 27
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27; Conser
                                                                         100.0%; Score 27; DB 10; Length 1729; llarity 100.0%; Pred. No. 0.0089; Conservative 0; Mismatches 0; Indels 0
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100.0%; Pred. No. 0.0
:ive 0; Mismatches
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Pred. No. 0
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0.0087;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 1350;
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                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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                                                                         Gaps
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                                                                         0;
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CURRENT APPLICATION NUMBER: US/09/822,827; CURRENT FILING DATE: 2001-03-28; NUMBER OF SEQ ID NOS: 982; SOFTWARE: FRASTSEQ for Windows Version 3.0; SEQ ID NO 980; LENGTH: 2406; TYPE: DNA; ORGANISM: Homo Sapiens
US-09-822-827-980
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.534C1
CURRENT APPLICATION NUMBER: US/09/822.827
CURRENT FILING DATE: 2001-03-28
NUMBER OF SEQ ID NOS: 982
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 977
LENGTH: 1758
TYPE: DNA
ORGANISM: Homo sapiens
US-09-822-827-977
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             οy
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US-09-822-827-980
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; Sequence 977, Application US/09822827

; Patent No. US20020081680A1
                                                                                                                                                                                                                                                                                                                                                                   RESULT 8
US-09-755-100-5
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APPLICANT: Xu, Jian
            GENERAL INFORMATION:
APPLICANY: SAVITZKY, Kinneret et al.
APPLICANY: SAVITZKY, Kinneret et al.
TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REFERENCE: 2786-0155P
CURRENT FILING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: IL 128587
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: IL 131363
PRIOR FILING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: IL 131363
PRIOR FILING DATE: 1999-08-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 100.0%; Score 27; Dest Local Similarity 100.0%; Pred. No. 0. Matches 27; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                Sequence 5, Application US/09755100 Patent No. US2002009189A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Xu, Jiangchun TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER FILE REFERENCE: 210121.534C1
NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1186 GTTCTGGTGCACCCCCAGTGGGTCCTC 1212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                778 GTTCTGGTGCACCCCAGTGGGTCCTC 804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           y Match 100.0%; Score 27; DB 10; Local Similarity 100.0%; Pred. No. 0.0091; hes 27; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 GTTCTGGTGCACCCCCAGTGGGTCCTC 27
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 Mismatches

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 2406;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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В
US-09-755-100-1
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US-09-755-100-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 9
US-09-755-100-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 5
LENGTH: 3423
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                  SOFTWARE:
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Application US/09755100 Patent No. US20020099189A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 100.0%; Score 27; DB 10; Best Local Similarity 100.0%; Pred. No. 0.0094; Matches 27; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Ver. SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: SAVITZKY, Kinneret et al.

TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES

FILE REFERENCE: 2786-0156P

CURRENT APPLICATION NUMBER: US/09/755,100:

CURRENT FILING DATE: 2001-01-08

PRIOR APPLICATION NUMBER: IL 128587

PRIOR FILING DATE: 1999-02-18

PRIOR FILING DATE: 1999-04-14

PRIOR FILING DATE: 1999-04-14

PRIOR FILING DATE: 1999-08-11

PRIOR FILING DATE: 1999-08-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 100.0%; Score 27; DB 10; Best Local Similarity 100.0%; Pred. No. 0.0094; Matches 27; Conservative 0; Mismatches 0;
                                                                                                                                           PRIOR APPLICATION NUMBER: IL 128587
PRIOR FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: IL 129439
PRIOR ETLING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: IL 131363
PRIOR FILING DATE: 1999-08-11
                                                                                                                                                                                                                                                                                                                         APPLICANT: SAVITZKY, Kinneret et al. TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES FILE REFERENCE: 2786-0156P
                                                                                                                                                                                                                                                                             CURRENT FILING DATE: 2001-01-08
                                                                                                                           NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 12
                 TYPE: DNA ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1658 GTTCTGGTGCACCCCCAGTGGGTCCTC 1684
                                                            LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 3846
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2081 GTTCTGGTGCACCCCCAGTGGGTCCTC 2107
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                                                              4661
                                                                                                       PatentIn Ver.
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Query Match

100.0%;

Score

27;

DB 10;

Length 4661;

Best Local Matches

27; Conservative (

0;

US-09-755-100-2

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Sequence 153, Application US/09925300
Patent No. US20020151681A1
GENERAL INFORMATION:
APPLICANT: Craig Rosen,
APPLICANT: Steve Ruben
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PA101
CURRENT APPLICATION NUMBER: US/09/925,300
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05988
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 2000-03-08
PRIOR FILING DATE: 1999-03-12
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 153
LENGTH: 1678
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Patent No. US/020099189A1

GENERAL INFORMATION:
APPLICAMT: SAVITERY, Kinneret et al.
TIFILE OF INVENTION: NOYEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REFERENCE: 2786-0156P
CURRENT FILING DATE: 12001-01-08
PRIOR APPLICATION NUMBER: US/09/755,100
CURRENT FILING DATE: 199-02-18
PRIOR APPLICATION NUMBER: IL 18587
PRIOR APPLICATION NUMBER: IL 18587
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-08-11
PRIOR FILING DATE: 1999-08-11
SOFTWARE: Patentin Ver. 2.1
SOFTWARE: Patentin Ver. 2.1
SOFTWARE: Patentin Ver. 2.1
LENGTH: 4661
                                                                                                                                                                               PEATURE:
NAME/REY: nisc feature
LOCATION: (1663)
OTHER INFORMATION: n equals a,t,g, or c
US-09-925-300-153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 100.0%; Score 27; DB 10; Length 4661; Best Local Similarity 100.0%; Pred. No. 0.0096; Matches 27; Conservative 0; Mismatches 0; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 GTTCTGGTGCACCCCAGTGGGTCCTC 27
                                                                      Score 23.8; DB 10;
Pred. No. 0.19;
D; Mismatches 2;
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; Mismatches 0;
                                                                                                                       Length 1678;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 101. Application US/09962832
Patent No. US20020110821A1
GENERAL INFORMATION:
APPLICANT: Ebner, Reinhard
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Usin
TITLE OF INVENTION: Sets
FILE REFERENCE: 689290-74
CURRENT APPLICATION NUMBER: US/09/962,832
CURRENT FILING DATE: 2001-09-25
PRIOR APPLICATION NUMBER: US/60/235,077
PRIOR RILING DATE: 2000-99-25
PRIOR RILING DATE: 2000-99-25
NUMBER OF SPO, ID NOS: 259
SOFTWARE: Patentin version 3.0
SEQ ID NO 101
LENGTH: 871
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Rosen et al.

FILE REFERENCE: PA105

FILE REFERENCE: PA105

CURRENT APPLICATION NUMBER: US/09/925,297

CURRENT FILING DATE: 2001-08-10

PRIOR APPLICATION NUMBER: PCT/US00/05989

PRIOR APPLICATION NUMBER: PCT/US00/05989

PRIOR APPLICATION NUMBER: 60/124,270

PRIOR APPLICATION NUMBER: 60/124,270

PRIOR APPLICATION NUMBER: 60/124,270

PRIOR APPLICATION NUMBER: 00/124,270

PRIOR APPLICATION NUMBER: US/09/925,297

PRIOR A
RESULT 15
US-09-880-107-2307
; Sequence 2307, Application US/09880107
; Patent No. US20020142981A1
; GENERAL INFORMATION:
; APPLICANT: Horne, Darci T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 13
US-09-925-297-286
""ence 286, Ap
                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-962-832-101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 14
US-09-962-832-101
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Best Local
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Patent No. US20020081659A1
                                                                                                                                                                                                                         194 TCCTGGTGCACCGCCAGTGGGTGCTC 219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       173 TCCTGGTGCACCGCCAGTGGGTGCTC 198
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                                                                                                                                                                                                                                                        2 TTCTGGTGCACCCCCAGTGGGTCCTC 27
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Local Similarity 88.5%; Pred. No. 2.3;
es 23; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                      Similarity 88.5
23; Conservative
                                                                                                                                                                                                                                                                                                                                                                  78.5%;
                                                                                                                                                                                                                                                                                                                                         Score 21.2; D
Pred. No. 2.3;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                              DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 10;
                                                                                                                                                                                                                                                                                                                                                                                       Length 871;
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                                                                                                                                                                                                                                                                                                                                      0,
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Sign

δÃ

-09-925-300-153

; TYPE: DNA; ORGANISM: Homo sapiens US-09-755-100-2

δÃ

Query Match Best Local : Matches

Local

Similarity 92.0 25; Conservative

88.1%; 92.6%;

0;

TYPE: DNA
ORGANISM: Homo sapiens

```
APPLICANT: Vockley, Joseph G.
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
FILE REFERENCE: 44921-5028-W0
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/231,379
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR FILING DATE: 2000-010-02
NUMBER OF SEQ ID NOS: 3950
SOFTWARR: Patentin Ver. 2.1
SEQ ID NO 2307
LENGTH: 871
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
OTHER INFORMATION: Genbank Accession No. US20020142981A1 M25629
US-09-880-107-2307
Ouery Match
PRIOR SIMILARITY 88.5%; Pred. No. 2.3;
Matches 23; Conservative 0; Mismatches 3; Indels 0; Gaps
O;
Matches 23; Conservative 0; Mismatches 3; Indels 0; Gaps
O;
Search completed: November 19, 2002, 10:16:26
Job time: 93.3846 secs
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Result
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Maximum DB seq
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Perfect score:
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              98765432
                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                         Score
           length: 0
length: .2000000000
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Match
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Gapop 10.0 , Gapext 1.0
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Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             | SIDS2/gcgdata/geneseq/geneseqn-embl/NA198B.DAT:*
| SIDS2/gcgdata/geneseq/geneseqn-embl/NA198B.DAT:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       \(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1980_DAT:*\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1981_DAT:*\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1982_DAT:*\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1983_DAT:*\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1983_DAT:*\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1984_DAT:*\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1985_DAT:*\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986_DAT:*\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986_DAT:*\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:*\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcgdata/geneseq/geneseq/geneseqn-embl/NA1986.DAT:\(\SIDS2\)/gcdata/geneseq/geneseq/geneseq/geneseq/geneseq/geneseq/geneseq/genese
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246
420
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492
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           118
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     AAD21363
AAT90073
AAT90074
ABV38513
ABV14335
ABV44545
ABV35419
ABV07463
ABV44248
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Prostate specific Prostate specific Prostate specific Prostate specific Human prostate exp Human prostate exp Human prostate exp Human prostate exp Human prostate exp
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22	23	23	23	23	23	23	23	23	24	16	17	21	19	20	18	19	23	24	22	22	23	23	21	23	23	20	18	21	23	24	20	23	23	18	23
F869	93	933	ABV29105	ABV29105	ABV24768	476	350	350	ABL69732	486	586	171	AAV32496	984	105	249	326	532	386	395	920	347	74	976	490	984	105	Z4713	507	620	94	813	22	29	ABV37397
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## ALIGNMENTS

RESULT 1
AAD21363
ID AAD2
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XX U 10-APR-2000; 2000US-195456P 10-APR-2001; 18-OCT-2001. WO200176622-A2 Unidentified. Prostate-specific antigen; PSA; immunogenic peptide; immune response; therapy; cancer; prostate cancer; tumour metastasis; prophylaxis; ds. Prostate specific antigen (PSA) derived peptide, CLP316 encoding DNA 28-JAN-2002 AAD21363; AAD21363 standard; (AVET ) AVENTIS PASTEUR LTD 2001WO-CA00473 (first entry) /\*tag= a
/product= "PSA derived peptide, CLP316"
/note= "CDS does not include start and stop codon" Location/Qualifiers 1..27 /partial DNA; 27 ВÞ

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Pedyczak A,

Chong

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CDY;

2001-663015/76.

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Prostate-specific antigen; PSA; epitope; vaccine; prostatic tumour cell; human leukocyte anticome; T lymbhoryte.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The patent discloses immunogenic peptides of prostate-specific antigen (PSA) and nucleic acids encoding them. The peptides of the invention are used to prepare a medicament to elicit an immune response in an animal. They are used to treat cancer such as prostate cancer and tumour metastasis. They are also useful for prophylaxis, for preparing monoclonal or polyclonal antibodies, and in conventional techniques of immunology, molecular biology, cell biology and recombinant DNA technology. The present sequence is a DNA encoding that the property of the present sequence is a DNA encoding that the present sequence is a DNA encoding tha
                                Prostate specific antigen multiple epitope peptide - useful processor of produce immune response against prostate specific in prostatic tumour cell
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New polypeptide, useful for comprises prostate-specific
                                                                                                                                                                                                                                                                                         Schlom J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-MAR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-MAR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Prostate specific antigen oligo-epitope encoding DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT90073 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PSA derived
                                                                                                                                                                                                                                                                                                                                                              (USSH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           lymphocyte;
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DB; AAW27387.
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27; Conserv
                                                                                                                                                                                                                                                                                                                                                          US SEC DEPT HEALTH.
US DEPT HEALTH & HUMAN SERVICES
                                                                                                                                                                                                                                                                                 Tsang K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 6 A; 6 C; 6 G; 9 T;
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peptide, CLP316.
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larity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    96US-0618936
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97WO-US04454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers 58..60
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                                                                                                                                                                                                                                                                                         Zaremba
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 27; DB 22;
Pred. No. 0.0051;
; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               prostate cancer,
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RESULT 3
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Best Local
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                                                                                                                                    Prostate specific antigen multiple epitope peptide - useful in vaccine to produce immune response against prostate specific antigen in prostatic tumour cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Prostate specific antigen; PSA; epitope; vaccine; immune respon prostatic tumour cell; human leukocyte antigen; HLA; cytotoxic; T lymphocyte; cancer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 90
                                                                                                                                                                                                                                  Schlom J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAT90074 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim
                                                                                                                                                                                                                                                            (USSH ) US SEC DEPT HEALTH.
(USSH ) US DEPT HEALTH & HUMAN SERVICES.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            l Similarity
27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               specific antigen oligo-epitope encoding DNA
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                                                                                                                                                                                                                                  Tsang K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CDNA;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immune response;
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The present sequence encodes a prostate specific antigen oligo-epitope peptide (PSA-OP) for use in a virux vector of the present invention. The PSA-OP comprises more than 1 adjoined PSA-OP which generates a prostate specific response in a portion of the human population. The PSA-OP can be used to produce PSA specific cytotoxic T lymphocytes; in vivo or in

can kill

prevent

growth of PSA

Claim 23; Page 67; 85pp; English

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RESULT 4

ARBV38513

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ID ABV38513

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products may be use
recombinant virus i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-FEB-2000;
16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
18-JUL-2000;
13-DEC-2000;
The invention relates to an isolated nucleic acid molecule (I) compa nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;

(b) monitoring the progression of prostate cancer in a patient;

(c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient;

(d) assessing the efficacy of a therapy for inhibiting prostate can patient;

(e) selecting a composition for inhibiting prostate cancer in a patient;
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                                                                                                                                                                                                                                                                                                                                                                                                                                WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; prostate pharmacogenomic
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                                                                                                                                                                                                                                                                                      Claim 1;
                                                                                                                                                                                                                                                                                                                               Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, uses for detecting presence of prostate cancer, stage of prostate cancer
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27; Conserv
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2000US-189862P.
2000US-207454P.
2000US-211314P.
2000US-219007P.
2000US-255281P.
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                                                                                                                                                                                                                                                                                      11750pp;
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Pred. No.
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                                                                                                                                                                                                                                                                                      English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         carcinogen;
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                                                                           prostate cancer
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                               patient;
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16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
18-JUL-2000;
                                                                                                                 The invention relates to an isolated nucleic acid molecule (I) com a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of t specification or its complement. (I) is useful for: (a) assessing whether a patient is afflicted with prostate cancer; (b) monitoring the progression of prostate cancer in a patient; (c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient.
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(h) asse
patient;
(I) is a
                                                                                      (C) assessing the efficacy of a therapy for
                                                                                                                                                                                                                         Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, use for detecting presence of prostate cancer, stage of prostate cancer
                    patient;
(I) is also useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; prostate pharmacogenomic

    (e) selecting a composition for inhibiting prostate cancer i
    (f) assessing the prostate cell carcinogenic potential of a
    (g) determining whether prostate cancer has metastasized in
    (h) assessing the aggressiveness or indolence of prostate ca

                                                                                     (d) assessing
in a patient;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    is also useful as a pharmacodyanamic or pharmacogenomic marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           determining whether prostate cancer has metastasized in a patient; assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAACTTCAGTGTGTGGACCTCCATGTT
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llarity 100.0%;
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marker;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      carcinogen;
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                                                                                                                                                                                                                                                                                                                    INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 23;
                     or,
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                     pharmacogenomic
                                                                                                inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pharmacodyanamic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                 prostate cancer
                                           cancer in
                                                      a patient
                                                                           in a patient;
                      marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                      patient;
                                                                                                                                                                    of the
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Length 429;

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RESULT 6
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                                 Query Match
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     Matches
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16-MAR-2000; 2000US-189862P.
25-MAY-2000; 2000US-207454P.
09-JUN-2000; 2000US-211314P.
18-JUL-2000; 2000US-255281P.
                                                                                                                                                        The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV062213) of the specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient;
(d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient;
(e) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient;
(f) assessing the prostate cell carcinogenic potential of a compound;
(f) assessing the prostate cell carcinogenic potential of a compound;
                                                                                                                                                                                                                                                                                                                                                                            Novel isolated nucleic acid molecule associated with cancerous state of prostate cells and correlating with presence of prostate cancer, useful for detecting presence of prostate cancer, stage of prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; prostate pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABV44545 standard; cDNA; 466
                                                                                                                                                                                                                                                                                                                                                   Claim 1; Page 8836; 11750pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Schlegel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200160860-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABV44545;
                                                                Sequence 466 BP; 98 A; 140 C; 136 G; 92 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MILL-) MILLENNIUM PREDICTIVE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-FEB-2001; 2001WO-US05171
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       selecting a composition for inhibiting prostate cancer in a patient; assessing the prostate cell carcinogenic potential of a compound; determining whether prostate cancer has metastasized in a patient, assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                               2001-662795/76.
                                                                                             also useful as a pharmacodyanamic or pharmacogenomic marker
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     l Similarity
27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           В,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Endege WO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cancer; cytostatic; carcinogen; pharmacodyanamic marker; marker; gene; ss.
                 100.0%;
     0;
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Score 27; DB 23;
Pred. No. 0.0086;
                                 Length 466;
     Indels
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   0;
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RESULT 8 ABV07463/c

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RESULT 7
ABV35419/c
ID ABV35419;
AC ABV35419;
XX AC ABV35419;
XX 16-SEP-2002 (first of the content of the conte
   В
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                                                                                                                Query Match
Best Local S
Matches 27
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16-MAR-2000; 2000US-1898629-
25-MAX-2000; 2000US-207454P.
09-JUN-2000; 2000US-211314P.
18-JUL-2000; 2000US-219007P.
13-DEC-2000; 2000US-255281P.
                                                                                                                                                                                                                                                                                                                              The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9, (ABM7000,10-ABM62213) of the specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient;
(d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient;
(e) assessing the prostate call carcinogenic potential of a compound;
(f) assessing the prostate cell carcinogenic potential of a compound;
(g) determining whether prostate cancer has metastasized in a patient;
(h) assessing the aggressiveness or indolence of prostate cancer in a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; prostate cancer;
pharmacogenomic marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 7378; 11750pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, uses for detecting presence of prostate cancer; stage of prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human prostate expression marker cDNA
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                                                                                                                                                                                                                                                                                    patient;(I) is also useful as a pharmacodyanamic or pharmacogenomic marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MILL-) MILLENNIUM PREDICTIVE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-FEB-2001; 2001WO-US05171.
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      286
ch 100.0%;
1 Similarity 100.0%;
27; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Endege WO,
                                                                                                                                                                                                                          103 A; 138 C; 136 G; 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytostatic; carcinogen; pharmacodyanamic gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Monahan JE;
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                                                                                                                0;
                                                                                                                Score 27; DB
Pred. No. 0.0
0; Mismatches
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                                                                                                                                             0.0086
                                                                                                                                                                                                                              T; 0 other;
                                                                                                                                                                          DB 23;
                                                                                                                                                                       Length 476;
                                                                                                                       Indels
                                                                                                                   0;
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useful
cer
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RESULT 9
ABV44248
ID ABV4
XX
AC ABV4
XX
DT 16-9
XX
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in a patient;
(e) selecting a com
(f) assessing the p
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(h) asse
patient;
(I) is a
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(a) assessing whether a patient is afflicted with prostate cancer;

(b) monitoring the progression of prostate cancer in a patient;

(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 1204-1205; 11750pp; English.
                                                                                                                                                                                                                                                                                                                                                            Novel isolated nucleic acid molecule associated with cancerous state of prostate cells and correlating with presence of prostate cancer, useful for detecting presence of prostate cancer, stage of prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-FEB-2000;
16-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; prostate pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human
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                                                                                                                                                                                                                                                                                                                                                                                                                    Schlegel
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09-JUN-2000;
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         16-SEP-2002
                           ABV44248;
                                             ABV44248 standard;
                                                                                                                                                                      Sequence 492
                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                      (MILL-) MILLENNIUM PREDICTIVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-JUL-2000;
13-DEC-2000;
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                                                                                            245
                                                                                                                                            Local
                                                                                                               Н
                                                                                                                                                                                                          selecting a composition for inhibiting prostate cancer in a patient; assessing the prostate cell carcinogenic potential of a compound; determining whether prostate cancer has metastasized in a patient; assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                               AAACTTCAGTGTGTGGACCTCCATGTT 27
                                                                                                                                                                                                                                                                                                                                                                                                  2001-662795/76
                                                                                            AAACTTCAGTGTGTGGACCTCCATGTT 219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               prostate
                                                                                                                                                                                          also
                                                                                                                                                                                                                                                                 in a patient;
                                                                                                                                  Similarity
27; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; 2000US-183319P.
; 2000US-189862P.
2000US-207454P.
; 2000US-211314P.
; 2000US-219007P.
; 2000US-255281P.
                                                                                                                                                                                          useful as
                                                                                                                                  Conservative
                                                                                                                                                                       BP; 102 A; 141 C; 115 G;
        (first
                                                                                                                                                                                                                                                                                                                                                                                                                    Endege
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expression marker cDNA 7454.
                                                                                                                                                                                                                                                        efficacy of a therapy
                                              CDNA;
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                                                                                                                                           100.0%;
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                                                                                                                                                                                         pharmacodyanamic or
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                                              616
                                                                                                                                                                                                                                                                                                                                                                                                                    Monahan JE;
                                                                                                                                  0
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                                              ВP
                                                                                                                                  Pred. No. 0. 
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDICINE INC
                                                                                                                                                     Score 27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             carcinogen;
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                                                                                                                                                                       134 T; 0 other;
                                                                                                                                                                                                                                                         for inhibiting prostate cancer
                                                                                                                                                     DB 23;
                                                                                                                                            .0086;
                                                                                                                                                                                        pharmacogenomic
                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pharmacodyanamic
                                                                                                                                                   Length 492;
                                                                                                                                  0
                                                                                                                                  Gaps
                                                                                                                                  0,
                                                                                                              RESULT 10
ABV37397/c
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Best Local S
Matches 27
                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                         cancer in a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAY-2000;
09-JUN-2000;
                 pharmacogenomic
                                                                16-SEP-2002
                                                                                   ABV37397;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schlegel R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pharmacogenomic
                                                                                                                                                                                                                                                                  (g) determining
(h) assessing
                                                                                                                                                                                                                                                                                                        in a patient;
                                                                                                                                                                                                                                                                                                                (d) assessing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-MAR-2000;
                                                                                                                                                     252
                                                                                                                                                                       Н
                                                                                                                                                                                                                                                is also useful as a pharmacodyanamic or pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001-662795/76
                                             prostate
                           prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   prostate
                                                                                                                                                                                         . Similarity
27; Conserv
                                                                                                                                                                                                                               616
                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                              BP;
                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Endege
                                                                                                                                                                                                                               122
                                                                                                                                                                                                                                                                                                                 efficacy of a therapy for inhibiting prostate cancer
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Human prostate expression marker cDNA 44239
                          (MILL-) MILLENNIUM PREDICTIVE MEDICINE
                                                         2000US-183319P.
2000US-189862P.
2000US-207454P.
2000US-211314P.
2000US-2119007P.
2000US-255281P.
                                                                                                                                                                   2001WO-US05171
                                                                                                                                                                                                                                                                                     cancer; cytostatic; carcinogen;
marker; gene; ss.
Monahan
JE;
                                                                                                                                                                                                                                                                                                                                                             ;•
_;
                                                                                             INC
                                                                                                                                                                                                                                                                                                       pharmacodyanamic marker;
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Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, used for detecting presence of prostate cancer, stage of prostate cancer tate of useful cer -

Claim 1; Page 8788; 11750pp; English.

The invention relates to an isolated nucleic acid molecule (I) com a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of t specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;

(b) monitoring the progression of prostate cancer in a patient;

(c) assessing the efficacy of a test compound to inhibit prostate ) comprising of the

(e) selecting a composition for inhibiting prostate cancer in a patient;(f) assessing the prostate cell carcinogenic potential of a compound;(g) determining whether prostate cancer has metastasized in a patient;

the aggressiveness or indolence of prostate cancer in a

marker

A; 176 Ç 186 G; 132 T; 0 other;

100.0%; Score 27; DB 23 Pred. No. 0.009; D; Mismatches 23; Length 616;

0;

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0;

Gaps

0

AAACTTCAGTGTGTGGACCTCCATGTT AAACTTCAGTGTGTGGACCTCCATGTT 278 27

ABV37397 standard; cDNA; 632

expression marker CDNA 37388

marker; gene; ss cancer; cytostatic; carcinogen; pharmacodyanamic marker;

Homo sapiens

WO200160860-A2.

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RESULT 11
AATT927
ID AATT92
XX XX AATT92
XX Z4-AF
DT 24-AF
XX CDNA
XX D1agr
KW D1agr
KW D056e
XX D006
XX HOMO
XX HOMO
XX HOMO
XX Y0973
XX 23-OC
XX 23-OC
                                                                                                                                                                                                                                                                В
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                                                                                                                                                                                                                                                                                                           Query Match 100.0%; Score 27; DB Best Local Similarity 100.0%; Pred. No. 0.00 Matches 27; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-FEB-2000;
16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
18-JUL-2000;
13-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient;
(d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient;
(e) assessing the prostate cancer in a patient;
(f) assessing the prostate cell carcinogenic prostate cancer in a patient;
(f) assessing the prostate cancer has metastasized in a patient;
(h) assessing the aggressiveness or indolence of prostate cancer in a patient;
                                                                                Diagnosis, prostate cancer; prostate specific antigen; PSA; metastatic-prostate cancer; prostate cell; secondary tumour; bone_metastatic anchoring; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel isolated nucleic acid molecule associated with cancerous state of prostate cells and correlating with presence of prostate cancer, useful for detecting presence of prostate cancer -
                                                                              metastatic-prost
                                                                                                                                cDNA for prostate specific antigen (PSA).
                                                                                                                                                                                                                                                                                                                                                             Sequence 632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 7678; 11750pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-662795/76.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-FEB-2001;
            23-OCT-1997
                                  WO9739139-A1
                                                                                                                                                         24-APR-1998
                                                                                                                                                                                                      AAT92976 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
                                                           Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                               pattent;
(I) is also useful as a pharmacodyanamic or pharmacogenomic marker.
                                                                                                                                                                                                                                                                326
                                                                                                                                                                                                                                                              70
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000US-207454P.
2000US-211314P.
2000US-219007P.
2000US-255281P.
                                                                                                                                                                                                                                                                                                                                                          BP; 130 A; 182 C; 160 G; 160 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000US-183319P
2000US-189862P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2001WO-US05171
                                                                                                                                                      (first entry)
                                                                                                                                                                                                        639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Monahan JE;
                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                300
                                                                                                                                                                                                                                                                                                                        DB 23; Length 632;
.009; ____
                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                  RESULT 12
ABV22297
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                   Schlegel
                                                                                                                                                                                                                                         Homo
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Query Match
                              17-FEB-2000; 2000US-183319P
16-MAR-2000; 2000US-189862P
25-MAY-2000; 2000US-207454P
09-JUN-2000; 2000US-211314P
18-JUL-2000; 2000US-21907P
13-DEC-2000; 2000US-255281P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CDNA sequences AAT92974-77 encode the prostate specific antigen (PSA). The detection of PSA mRNA in peripheral blood is associated with metastatic prostate cancer. Detection of PSA mRNA in the bloodstream indicates that prostate cells are circulating in the blood and confirms diagnosis of cancer. Circulating prostate cells also indicate a risk of a secondary tumour or bone metastatic anchoring. Detection of the PSA mRNA after prostatectomy or radiotherapy indicates a risk that the prostate cancer has spread and that the surgery and radiotherapy was not effective. PCR primers AAT92971-73 were used for the monitoring and diagnosis of patients with prostate cancer. The method comprises the detection of PSA mRNA by reverse transcriptase polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                  Human prostate expression marker cDNA 22288
                                                                                                                                                                                                                                                                                                                                                                    13-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                  ABV22297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 16; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-APR-1997;
                                                                                                                                                                                    23-AUG-2001.
                                                                                                                                                                                                                   WO200160860-A2
                                                                                                                                                                                                                                                                                  Human; prostate pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                                                                     ABV22297;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Monitoring or diagnosis of prostate oppositate-specific antigen mRNA using
(MILL-)
                                                                                                                                                   20-FEB-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RT-PCR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (SMIK ) SMITHKLINE BEECHAM CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  486 AAACTTCAGTGTGTGGACCTCCATGTT
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                                                                                                                                                                                                                                                 sapiens.
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27; Conserv
 MILLENNIUM PREDICTIVE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                  standard; cDNA; 651
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP;
                                                                                                                                                   2001WO-US05171
                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                cancer; cytostatic; carcinogen;
marker; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         133 A; 191 C; 185 G; 130 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    512
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g specific
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0.0091;
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primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                   pharmacodyanamic marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 639;
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THE REAL STATES OF THE PROPERTY OF THE PROPERT
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ABV28130
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:
(a) assessing whether a patient is affiliated with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                                                                                                                                                                                     17-FEB-2000;
16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
18-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; prostate pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (c) assessing the efficacy of a test compound to inhibit cancer in a patient;(d) assessing the efficacy of a therapy for inhibiting in a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel isolated nucleic acid molecule associated with cancerous state of prostate cells and correlating with presence of prostate cancer, useful for detecting presence of prostate cancer, stage of prostate cancer.
                                                                                                                                                          WPI;
                     Claim
                                                                   ior
                                                                                       prostate
                                                                                                            Novel isolated
                                                                                                                                                                                                        Schlegel
                                                                                                                                                                                                                                                    (MILL-)
                                                                                                                                                                                                                                                                                                  13-DEC-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-SEP-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABV28130 standard; cDNA; 651
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              £9£0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             invention relates
                                                              detecting
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|||||||||||||||
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sapiens
                                                                                                                                                          2001-662795/76
                     1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      also useful as a pharmacodyanamic or pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
27; Conserv
                                                                                                                                                                                                                                                    MILLENNIUM PREDICTIVE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  651
                                                                                                                                                                                                      70
                   Page
                                                                                         cells
                                                                                                                                                                                                                                                                                             ; 2000US-183319P.
; 2000US-189862P.
; 2000US-207454P.
; 2000US-211314P.
; 2000US-219007P.
; 2000US-255281P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               2001WO-US05171.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
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                                                              presence of prostate
                     5827;
                                                              nucleic acid molecule associated with cancerous state
and correlating with presence of prostate cancer, use
presence of prostate cancer, stage of prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      expression marker cDNA 28121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer;
marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             164 A; 171 C;
                   11750pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%;
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given
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cytostatic;
                                                                                                                                                                                                        Monahan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    isolated nucleic acid molecule (I) comprising in Tables 1-9 (ABV00010-ABV62213) of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 27;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               156
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        carcinogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               160 T; 0 other;
                                                                                                                                                                                                                                                      INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 651;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          marker;
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RESULT 14
AAX08947
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                     (KLEE/)
(MAYO-)
(TIND/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to an isolated nucleic acid molecule (I) com a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of t specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;

(b) monitoring the progression of prostate cancer in a patient;

(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                              WPI;
                                                                                                                                     Klee
                                                                                                                                                                                                                                                                                                                                  CDS
                                                                                                                                                                                                                                                                                                                                                                     Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAX08947;
                                                                  the
                                                                                                  P-PSDB;
                                                                                                                                                                                                                                                                                      WO9859073-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                            Human prostate specific antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX08947
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer in a patient;
(d) assessing the efficacy of a therapy for
                                           Disclosure;
                                                                                                                                                                                                                  20-JUN-1997;
                                                                                                                                                                                                                                         19-JUN-1998;
                                                                                                                                                                                                                                                                30-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                          Human kallikrein 2; hK2; breast cancer; diagnosis; detecti
treatment; monitoring; prostate specific antigen; PSA; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            in a patient;
                                                                                                                                                           ( /NDOA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        249
                                                               of human kallikrein 2\, - as a marker for developing products for diagnosis, prognosis, monitoring and treatment of breast cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       is also useful as a pharmacodyanamic or pharmacogenomic marker
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B
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                                                                                                             1999-105632/09
                                                                                                                                                      ) KLEE G G.
) MAYO FOUNDATION.
) TINDALL D J.
) YOUNG C Y F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27;
                                                                                                    AAW96187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  651
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard;
                                                                                                                                     Tindall DJ,
                                          Page 34-35; 67pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                    97US-0050963
                                                                                                                                                                                                                                         98WO-US12840
                                                                                                                                                                                                                                                                                                             /label=
                                                                                                                                                                                                                                                                                                                         /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 164 A; 171 C;
                                                                                                                                                                                                                                                                                                                                              ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CDNA;_711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                   Young
                                                                                                                                                                                                                                                                                                    a
Prostate_specific_antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                     CYF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 27;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                             (PSA) gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .0091;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
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                                                                                                                                                                                                                                                                                                                                                                                                       detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            651;
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Human kallikrein 2 (hK2) is expressed at elevated levels relative the prostate cancer antigen, prostate specific antigen (PSA) by breast cancer cells when stimulated by androgens. Detecting level

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RESULT 15
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Best Local S
Matches 27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; prostate specific antigen; PSA; cytostatic; antiviral; immunostimulant; cell-mediated immune response; tumour; breast cancer; virus infection; prostate cancer; colorectal cancer; pancreatic cancer; lymphoma; leukaemia; hepadnavirus; lentivirus; herpesvirus; human immunodeficiency virus; HIV; flavivirus; pestivirus; gene; ss.
   Inducing a cell-mediated immune response against a target antigen, reducing undesired cells and stimulating presentation of an antigen a cell, comprises administering a polynucleotide encoding a variant an antigen -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 711 BP; 148 A; 208 C; 206 G; 149 T; 0 other;
an antigen
                                                                                                                                                                                                                                                                                 (AMBI-) AMERICAN FOUND BIOLOGICAL RES (MINC) MINCHEF M S. (LOUK/) LOUKINGV D I. (ZOUB/) ZOUBAK S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-NOV-2001; 2001WO-US45626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200240059-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cDNA encoding human prostate specific antigen (PSA) variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-SEP-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABK86205
                                                                                                                                          P-PSDB;
                                                                                                                                                                                                                              Mincheff MS,
                                                                                                                                                                                                                                                                                                                                                                                                                               01-NOV-2000; 2000US-0704232.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sapiens.
                                                                                                                                          2002-527524/56.
DB; AAU98921.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l Similarity
27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; cDNA; 744
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            llarity 100.0%; Score 27; DB 20; Length 711; 100.0%; Pred. No. 0.0092; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                              Loukinov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
7..744
/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /product= "Prostate specific antigen (PSA) variant
    with signal sequence deleted"
                                                                                                                                                                                                                              Zoubak S
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by
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The invention

8

a method of inducing

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cell-mediated

Disclosure; Page 121-122; 146pp; English.

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response against a cell comprising a target antigen (I) in a subject, treating a subject having undesired cells, for example tumour cells or virally infected cells (C), reducing the number of (C) in a subject and stimulating presentation of (I) by a cell. This is done by administering a polynucleotide (II) encoding a variant of (I), so that (II) expressed in a cell and cell-mediated immune response is induced. The method can be used to treat prostate cancer, breast cancer, colorectal cancer and pancreatic cancer, as well as lymphomas and leukaemias. The method is also useful in treating chronic viral infections such as those caused by hepadnaviruses, lentiviruses (including human immunodeficiency virus (HIV)), herpesviruses and the slaviviruses and pestiviruses. The present sequence represents the cod sequence of human prostate specific antigen (PSA) variant which has the signal sequence deleted, used as a target antigen in the method of the invention
                                                   of the invention.
                                                                                    which
in the method
                                                                                                                                                                                                                                                                                                                                                                                       a subject,
                                                                                                                                                                                                                                                                                                                                   so that
                                                                                                                                    coding
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Sequence 744 BP; 153 A; 222 C; 213 G; 156 T; 0 other;

밁 QY Query Match
Best Local Similarity
Matches 27; Conserv 466 1 ANACTTCAGTGTGTGGACCTCCATGTT 27 AAACTTCAGTGTGTGGACCTCCATGTT 492 100.0%; 9 ilarity 100.0%; For Conservative 0; Score 27; Pred. No. Mismatches DB 24; Length 744; 0.0093; hes 0; Indels 0, Gaps

0;

2002, 03:06:38

Search completed: November 19, Job time: 191.769 secs

GenCore version Copyright (c) 1993 - 2002

Compugen Ltd 5,1.3

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Result
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Minimum
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Title:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score
     nucleic search, using sw model
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1: /cgn2_6/ptodata/2,
2: /cgn2_6/ptodata/2,
3: /cgn2_6/ptodata/2,
4: /cgn2_6/ptodata/2,
5: /cgn2_6/ptodata/2,
6: /cgn2_6/ptodata/2,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          length: 0
length: 2000000000
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27
1 aaacttcagtgtgtggacctccatgtt 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          441362 seqs, 153338381 residues
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /cgn2_6/ptodata/2/ina/5A_COMB.seq:*
/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
/cgn2_6/ptodata/2/ina/6A_COMB.seq:*
/cgn2_6/ptodata/2/ina/6B_COMB.seq:*
/cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/2/ina/backfiles1.seq:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
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                                             US-09-100-264-4
US-08-358-820-13
US-08-605-785-616
US-08-605-785-616
US-08-64-527A-14
US-08-64-527A-14
US-08-64-527A-14
US-08-64-527A-1
US-08-56-105-31
US-08-56-993-1
US-08-56-993-1
US-08-56-993-7
US-08-622-046B-2
US-08-767-829A-7
US-08-767-829A-7
US-08-767-829A-7
US-08-768-859A-7
US-08-768-859A-7
US-08-768-859A-9
US-08-622-046B-13
US-08-622-046B-6
US-08-622-046B-6
US-08-622-046B-6
US-08-622-046B-6
US-08-622-046B-6
US-08-622-046B-6
US-08-622-046B-6
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156.345 Million cell updates/sec
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13, Appl
616, App
14, Appl
14, Appl
14, Appl
15, Appli
17, Appli
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28	18.2	67.4	832	ω	US-08-768-859A-20	Sequence 20, Appl
29	18.2	67.4	832	w	US-08-767-820A-5	~
30	18.2	67.4	832	w	US-08-767-820A-20	20
31	18.2	67.4	832	ω	US-08-622-046B-4	Sequence 4, Appli
32	18.2	67.4	832	ω	US-08-622-046B-15	
ယ	18.2	67.4	832	Ŋ	PCT-US95-06157-5	Sequence 5, Appli
34	18.2	67.4	1341	4	US-08-983-075D-6	Sequence 6, Appli
35	18.2	67.4	1358	4	US-08-983-075D-8	Sequence 8, Appli
36	18.2	67.4	2417	4	US-09-605-785-334	
37	18.2	67.4	2417	4	US-09-439-313-334	Sequence 334, App
38	18.2	67.4	2417	4	US-09-352-616A-334	Sequence 334, App
39	18.2	67.4	2417	4	US-09-232-149A-334	Sequence 334, App
40	18.2	67.4	3674	4	US-09-605-785-698	Sequence 698, App
41	18	66.7	3360	_	US-08-408-093-5	Sequence 5, Appli
42	18	66.7	3360	۲	US-08-408-420A-5	
43	18	66.7	3360	_	US-08-714-901-5	
44	18	66.7	3360	w	US-08-040-741-5	Sequence 5, Appli
c 45	17.2	63.7	1548	4	US-09-099-053-1	•
					ALIGNMENTS	
RESULT 1						

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GENERAL INFORMATION:
APPLICANT: COCCE, Carlo
APPLICANT: Gomella, Leonard
APPLICANT: Mulholland, S. Grant
APPLICANT: Mulholland, S. Grant
APPLICANT: Moreno, Jose
APPLICANT: Fischer, Rainer
TITLE OF INVENTION: Methods of Detecting Micrometa:
NUMBER OF SEQUENCES: 14
CORRESSEDNUBENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz &
STREET: One Liberty Place 46th. Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-100-264-4; Sequence 4, Applica; Patent No. 6235486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 711
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-100-264-4
                                                                                                                                                                                                                                                                                                                                                                    RESULT 2
US-08-358-782D-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 27
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APPLICANT: "Young" Charles Y.F.
APPLICANT: Tindall, Donald J.
APPLICANT: Klee, George G.

TITLE OF INVENTION: Method for detection of breast cancer
FILE REFERENCE: 545.006US1
CURRENT APPLICATION NUMBER: US/09/100,264
CURRENT FILING DATE: 1998-06-19
EARLIER APPLICATION NUMBER: US 60/050,963
EARLIER FILING DATE: 1997-06-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 4
LENGTH: 711
                                                                                                                                                                                                                                                                                                                         Sequence 13, Application US/08358782D Patent No. 5674682
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS:
COMPUTER READABLE FORM
               STREET: One Liberty CITY: Philadelphia STATE: PA ZIP: 19103
                                                                                                                                                                                                                                                                                                                                                                                                                                                        436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 AAACTTCAGTGTGTGGACCTCCATGTT 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAACTTCAGTGTGTGGACCTCCATGTT 462
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27; Conserv
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                                                                                                                                                                        Micrometastasis
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                                                                                                                                                                        of Prostate
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MEDIUM TYPE: COMPUTER: II

: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS

OPERATING SYSTEM:

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MOLECULE TYPE: DNA (genomic);
HYPOTHETICAL: NO
ANTI-SENSE: NO
US-08-358-782D-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-764-527A-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
TELEPAX: 215-568-3439
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 992 base pairs
TYPE: nucleic acid
STRANDEDNESS: Single
TOPOLOGY: 11 near
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 100.
Best Local Similarity 100.
Matches 27; Conservative
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/358,782
FILING DATE: 15-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: BEATGELL, LOTI Y.
REGISTRATION NUMBER: 34,293
REFERENCE/DOCKET NUMBER: TJU-1327
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
COMPATION WIMPERS. II CAN JOE 1677
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Gomella, Leonard
APPLICANT: Mulholland, S. Grant
APPLICANT: Moreno, Jose
APPLICANT: Fischer, Rainer
TITLE OF INVENTION: Methods of Detecting Micrometastasis
TITLE OF INVENTION: Prostate
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Beardell, Lori Y.
REGISTRATION NUMBER: 34,293
REFERENCE/DOCKET NUMBER: TJU-1327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Re
CURRENT APPLICATION DATA
APPLICATION NUMBER: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  551 AAACTTCAGTGTGTGGACCTCCATGTT 577
                                                                                                                                                                                                                                                                                                                                                                                            STREET: One Liberty
CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                          APPLICATION NUMBER: US/08/764,527A FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 AAACTTCAGTGTGTGGACCTCCATGTT 27
                                                                                                                                                                       ASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13, Application US/08764527A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NFORMATION:
                                                                                                                                                                                                                                                                                                                                                               19103
                                                                                                                                                                                                                                                                                                                                                                                                                      E: Woodcock Washburn Kurtz Mackiewicz & One Liberty Place 46th. Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-DEC-1994
N: 435
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wang, Aijun
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C16
CURRENT APPLICATION NUMBER: US/99/605,785
CURRENT FILING DATE: 2000-06-27
NUMBER OF SEQ ID NOS: 835
SOFTWARE: FASTSEQ FOR Windows Version*3.0
SEQ ID NO 616
SEQ ID NO 616
SEQ ID NO 616
LENGTH: 1350
TYPE: DNA
ORGANISM: Homo sapien
US-09-605-785-616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; HYPOTHETICAL: II; ANTI-SENSE: NO US-08-764-527A-13
                                                                                                        US-08-358-782D-14
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                                                          Patent No.
GENERAL /
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 100.0%; Score 27; DB 2; Best Local Similarity 100.0%; Pred. No. 0.00114 Matches 27; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
Patent N
                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                               MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 551 AAACTTCAGTGTGGACCTCCATGTT 577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 AAACTTCAGTGTGTGGACCTCCATGTT 27
                                                                                                                                                                                                                          Local Similarity es 27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENGTH:
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                                                       NEORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleic acid
                                                                                        Application US/08358782D
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mitcham, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Henderson,
                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              iang, Yuqui
Gomella, Leonard
Mulholland, S. Grant
Moreno, Jose
                                           Croce,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA (genomic)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Robert A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13:
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                                                                                                                                                                                                                                                                                                                                      Sequence 1
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                          GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS.
                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: 215-568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Beardell, Lori Y.
NAME: Beardell, Lori Y.
A 293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HYPOTHETICAL:
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: DN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Rel
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
               CURRENT APPLICATION DATA:
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                                                                                               STATE: 19103
                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                  TITLE OF INVENTION: Methods of Detecting Micrometastasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                               532 AAACTTCAGTGTGTGGACCTCCATGTT 558
                                                                                                                                                                                                                                                                 PPLICANT:
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FILING DATE: 15-DEC
CLASSIFICATION: 435
                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 34
                               SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM:
 APPLICATION NUMBER:
                                                                                                                                                                      ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                      14,
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                                                                                                                                                                                                                                                                                                                          INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nucleic acid
EDNESS: single
                                                                                                                                      Philadelphia
                                                                                                                                                                                                                                                                                                                                                      Application
                                                                                                                                                 E: Woodcock Washburn Kurtz Mackiewicz & One Liberty Place 46th. Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1462 base pairs
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                                                                                                                                                                                                                                                                            Mulholland, S. Grant
                                                                                                                                                                                                                                                              Moreno, Jose
                                                                                                                                                                                                                                                                                                            Croce
                            PatentIn Release #1.0, Version #1.25
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SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                 Prostate
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                                                                                                                                                                                                                                                 Rainer
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                                                                                                                                                                                                                                                                                          Leonard
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US/08/764,527A
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US-08-764-527A-14
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Best Local Similarity
Matches 27; Conserv
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                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE NU HYPOTHETICAL: NU
                                                                                                                                                                        APPLICATION NUMBER: 08/2 FILING DATE: 15-APR-1994 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                              TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
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LENGTH: 1462 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: single
                                                                                                                                             NAME: White, John P
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                APPLICATION NUMBER: FILING DATE: 18-APP
                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
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                                                                                   TELEFAX:
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REGISTRATION NUMBER:
                           LENGTH:
                                                                                                  TELEPHONE:
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E: New York
TRY: U.S.A.
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          nucleic acid
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                                                                     ONE: (212) 977-95
422523 COOP UI
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                                                                                                              John P.
John P.
NUMBER: 28,678
NUMBER: 43677/JPW/TEP
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                                                                                                                                                                                                                                                                                          Version #1.25
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Best Loc
Matches
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                                                                                                                Matches
                                                                                                                                                     Query Match
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Best Local Similarity
Matches 27; Conser
                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS: LENGTH: 1729 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 4367
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 278-0400
TELEFAX: (212) 391-0525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/718,547
FILING DATE:
CLASSIFICATION: 10.0435
APPLOADEY ACCESS 10.045
                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
ORIGINAL SOURCE:
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TITLE OF INVENTION: A MG
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                                       813 AAACTTCAGTGTGTGGACCTCCATGTT 839
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LOCATION:
                                                                                                                                                                                                                                 NAME/KEY:
                                                                                                                                                                                                                                                                        ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: UZIP: 10036
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STATE: New York
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                                                            1 AAACTTCAGTGTGTGGACCTCCATGTT 27
                                                                                                              Local Similarity es 27; Conser
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1185 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          U.S.A.
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                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Katz, Aaron E.; Buttyan, Ralph; Raffo, Anthony;
                                                                                                                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                      Homo sapiens
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                                                                                                                                100.0%;
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                                                                                                                                  Score 27;
Pred. No.
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US-08-294-611-1/c
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US-09-056-105-31
; Sequence 31, A
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APPLICANT: KIPPS, THOMAS J.

APPLICANT: WU, YUNQI
TITLE OF INVENTION: VACCINES WITH ENHANCED INTRACELLULAR
TITLE OF INVENTION: PROCESSING
FILE REFERENCE: 233/221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, Application US/08294611 Patent No. 5506106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/056,105 CURRENT FILING DATE: 1998-04-06 EARLIER APPLICATION NUMBER: 60/043,467 EARLIER FILING DATE: 1997-04-10 NUMBER OF SEQ ID NOS: 35
                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/2
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                           TELEPHONE: (215) 568-3100
TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4200 AAACTTCAGTGTGTGGACCTCCATGTT 4226
                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: LOTIY: Beardell
REGISTRATION NUMBER: 34,293
REFERENCE/DOCKET NUMBER: TJ
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: U
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Methods of DetecTITLE OF INVENTION: Micrometastasis NUMBER OF SEQUENCES: 2
                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: One Liberty
CITY: Philadelphia
STATE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Woodcock Washburn Kurtz
ADDRESSEE: Mackiewicz & No. 5506106ris
STREET: One Liberty Place - 46th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 AAACTTCAGTGTGTGGACCTCCATGTT 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity es 27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1, Application US/09056105 6287569
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FastSEQ for Windows Version 3.0
Nucleic
EDNESS: Single
GY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; ilarity 100.0%; Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Methods of Detecting .
Micrometastasis Of Prostate Cancer
                                                                                                                                                                                                                                                                        US/07/973,322
                                                                                                                                                                                                                                                                                                                                                    US/08/294,611
                                                                                                                                                                         TJU-0722
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Pred. No. 0.0018;
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Mismatches

Length 7130; Indels

0;

0;

Mulholland, S.

Grant

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RESULT 12
US-08-596-993-1/c
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US-08-358-782D-1/c
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US-08-294-611-1
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                                          Sequence 1, Application Patent No. 5688649
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                                                                                                                                                                              Matches
                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
GENERAL INFORMATION:
APPLICANT: Croce, Carlo
APPLICANT: Gomella, Leonard
                                                                                                                                                                                                                                                                                                                                           TELEFAX: 215-568-3439
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/358
FILING DATE: 15-DEC-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Beardell, Lori Y.
REGISTRATION NUMBER: 34,293
REFERENCE/DOCKET NUMBER: TJU-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Croce, Carlo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                  MOLECULE TYPE: DI
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Fischer, Rainer
TITLE OF INVENTION: Methods of Detecting Micrometastasis of Prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
                                                                                                                           20
                                                                                                                                                                                                                                                                                                   STRANDEDNESS:
                                                                                                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE:
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                                                                                                                                       2 AACTTCAGTGTGTGGACCTC 21
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                                                                                                                                                                                                                                                                                                                               ENGTH:
                                                                                                                         AACTTCAGTGTGGACCTC 1
                                                                                                                                                                               20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Philadelphia
                                                                                                                                                                                                                                                                                                                 nucleic acid
                                                        Application US/08596993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Application US/08358782D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    E: Woodcock Washburn Kurtz Mackiewicz & One Liberty Place 46th. Floor
                                                                                                                                                                                                                                                                                                                               20 base pairs
                                                                                                                                                                                                                                                                                                                                                                       215-568-3100
215-568-3439
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                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                           linear
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                                                                                                                                                                                             Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                 TJU-1327
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                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                        1;
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Query Match
Best Local Similarity
Vatches 20; Conserv?
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US-08-764-527A-1/c
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US-08-596-993-1
                                                                                                                                                                                                                                                          Sequence 1
Patent No.
                                                                                               APPLICANT: Croce, Carlo
APPLICANT: Gomella, Leonard
APPLICANT: Mulholland, S. Grant
APPLICANT: Moreno, Jose
APPLICANT: Fischer, Rainer
TITLE OF INVENTION: Methods of De
TITLE OF INVENTION: Prostate
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/294,611
FILING DATE: 23-AUG-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/973,322
FILING DATE: 29-OCT-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (215) 568-34
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Mulholland
APPLICANT: Moreno, Jo
APPLICANT: Fischer, R
TITLE OF INVENTION: M
TITLE OF INVENTION: M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: (215) 568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
          STREET: ONE L.CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                      CORRESPONDENCE ADDRESS:
STATE: F...
19103
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: One LINELLY CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/596,993 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: UZIP: 19103
                                                                                                                                                                                                                                                                                                                                                             20 AACTTCAGTGTGTGGACCTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Mackiewicz
                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                           AACTTCAGTGTGTGGACCTC 21
                                                                                                                                                                                                                                                          1, Application US/08764527A
5. 5939258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mark DeLuca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PA
                                                       E: Woodcock Washburn Kurtz Mackiewicz
One Liberty Place 46th. Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E: Mackiewicz & No. 5688649ris
One Liberty Place - 46th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
                                                                                                                                                                                                                                                                                                                                                                                                                            74.1%; Score 20; llarity 100.0%; Pred. No. Conservative 0; Mismatcl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Moreno, Jose G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IBM COMPATIBLE SYSTEM: PC-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Woodcock Washburn Kurtz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DISKETTE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                568-3439
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Methods of Detecting
Micrometastasis Of Prostate Cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TJU-1657
                                                                                                                                         Detecting
                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                           Micrometastasis
                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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STREET: D.O. Box 2938

CITY: Minneapolis

STATE: M.

COUNTRY: USA

ZIE: 55402

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: EBM PC compartable

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFFWARE: MS-LCATION DATA:

APPLICATION NUMBER: US/08/622,046B

CLASSIFICATION: 514

ATTORNEY/AGENT INFORMATION:

REGISTRATION NUMBER: 30,440

REGISTRATION NUMBER: 30,440

REGISTRATION NUMBER: 30,440

REFERENCE/DOCKET NUMBER: 30,440

TELEPHONE: 612-373-6903

TELEPHONE: 612-373-6903

TELEPHONE: 612-373-6903

TELEPHONE: 612-373-9330

TELEPHONE: 612-373-393-3061
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION UNMBER: US/08/358.782
FILING DATE: 11-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: Beardell, Lori Y.
REGISTRATION UNMBER: 34,293
REFERENCE/DOCKET NUMBER: 7JU-1327
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDENNESS: single
TOPOLOGY: linear
TOPOLOGY: linear
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 14
US-08-622-046B-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: DNA
HYPOTHETICAL: NO
HATT-SENSE: NO
US-08-764-527A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 2, Application US/08622046B Patent No. 6103237
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ouery Match 74.1%; Score 20; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 1.2; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Saedi, Mohamm
APPLICANT: Mikolajczyk,
TITLE OF INVENTION: Stab
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth, P.A. STREET: P.O. Box 2938
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mohammed S
                                                                                                                                                                                                                                                                                                                                                                                                                       zyk, Stephen D.
Stable Variant HK2 Polypeptide
21
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                            Search completed: November 19, 2002, 05:20:22 Job time: 53:9615 secs
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COMPUTER READMALE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
EVERNIT APPLICATION DATA:
APPLICATION NUMBER: US/08/622.046B
FILING DATE: 26-MARCH-1996
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: WOSSINGT, Warren D.
REGISTRATION NUMBER: 30.40
REGISTRATION UNDER: 476.001US1
TELEPAN: 612-373-6903
INFORMATION FOR SED ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 711 base pairs
TYPE: nucleic acid
STRANDENESS: double
TOPOLOGY: linear
MOLECULE TYPE: CDNA
                                                                     Query Match
Best Local S
Matches 20
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; TYPE: nucleic acii;
; STRANDEDNISS: doi:
; TOPOLOGY: linear
; MOLECULE TYPE: CDNA
US-08-622-046B-2
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US-08-622-046B-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 13, Application US/08622046B
Patent NO. 6103237
GENERAL INFORMATION:
APPLICANT: Saedi, Mohammed S.
APPLICANT: Mixolajczyk, Stephen D.
TITLE OF INVENTION: Stable Variant HK2-Polypeptide
NUMBER OF SEQUENCES: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 67.4%;
Best Local Similarity 87.0%;
Matches 20; Conservative
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth, P.A.
STREET: P.O. Box 2938
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STALL COUNTRY: 05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Schweg
STREET: P.O. Box
CITY: Minneapolis
STATE: MN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4 CTTCAGTGTGTGGACCTCCATGT 26
                                                              Similarity 87.0
20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      double
                                                                             67.4%;
87.0%;
                                                                0;
                                                                             Score 18.2;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 18.2; DE Pred. No. 14; 0; Mismatches.
                                                            Mismatches
                                                                                          DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 3;
                                                                                        ω
••
                                                                                      Length 711;
                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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